

TAB]

AGI code	Agrisera antibody name	Product number
AT1G02280	Toc33 chloroplast outer envelope membrane translocon complex protein	AS07 236
AT1G03130	PsaD PSI-D subunit of photosystem I	AS09 461
AT1G06680	PsbP 23 kDa protein of the oxygen evolving complex (OEC) of PSII	AS06 167
AT1G06760.1	H1 histone H1	AS11 1801
AT1G07510	FtsH10 plant fts metallo-protease H10	AS07 251
AT1G07940	EF1A elongation factor 1-alpha / EF-1-alpha	AS11 1633
AT1G08380	PsaO PSI-O subunit of photosystem I	AS04 050
AT1G09340	CSP41b ribosome associated endonuclease (CRB)	AS08 298
AT1G09570	Phytochrome A photoreceptor PhyA protein	AS07 220
AT1G09640	eEF1B-gamma1 and 2 elongation factor 1-gamma 1 and 2	AS10 676
AT1G09700	HYL1 hyponastic leave phenotype ds-RNA binding protein	AS06 136
AT1G09940	GluTR glutamyl -tRNA reductase	AS10 689
AT1G11510	DNA-binding storekeeper protein-related transcriptional regulator	AS11 1774
AT1G12030	HSP17.7 cytosolic class II heat shock protein 17.7	AS07 255
AT1G12520	CCS chloroplastic copper chaperone for SOD	AS07 219
AT1G15690	V-PPase vacuolar H ⁺ -pyrophosphatase	AS09 465
AT1G15820	Lhcb6 CP24 chlorophyll a/b-binding protein of plant PSII	AS01 010
AT1G19570	DHAR1 Dehydroascorbate Reductase 1	AS11 1746
AT1G20340	PC plastocyanin	AS06 141
AT1G22840	Cyt c cytochrome c	AS08 343
AT1G23490	AtArf2 ADP ribosylation factor 2	AS09 476
AT1G28320	DEG15 endopeptidase, peroxisomal marker	AS10 711
AT1G29910	Lhcb1 LHCII type I chlorophyll a/b-binding protein (100 µl)	AS01 004
AT1G30230	eEF1B-beta1 and 2 elongation factor 1-beta1 and 1-beta2	AS10 677
AT1G30380	PsaK PSI-K subunit of photosystem I	AS04 049
AT1G30400	AtMRP1, AtABCC1 ABC transporter C family member 1	AS09 472
AT1G31330	PsaF PSI-F subunit of photosystem I	AS06 104
AT1G31810	AtFH14 Formin-like protein 14 (50 ug)	AS11 1750
AT1G32060	PRK ribulose-5-P-kinase phosphoribulokinase	AS07 257
AT1G43670	cFBPase cytosolic fructose-1,6-bisphosphatase (cytoplasm marker in photosynthetic tissue)	AS04 043
AT1G44575	PsbS 22 kDa Lhc-like PSII protein	AS09 533
AT1G45474	Lhca5 PSI type V chlorophyll a/b-binding protein	AS05 082
AT1G48410	AGO1 argonaute 1	AS09 527

TAB]

AT1G53540	HSP17.6 cytosolic class I heat shock protein 17.6	AS08 284
AT1G54780	Tlp18.3 Thylakoid lumen 18.3 kDa protein	AS08 369
AT1G55670	PsaG PSI-G subunit of photosystem I	AS04 048
AT1G59870	AtPDR8 ABC transporter G family member 36	AS09 471
AT1G61520	Lhca3 PSI type III chlorophyll a/b-binding protein	AS01 007
AT1G62300	AtWRKY6 WRKY transcription factor 6	AS11 1778
AT1G62360	STM homeobox protein SHOOT MERISTEMLESS	AS11 1764
AT1G62480	AtCCaP1 vacuolar calcium-binding protein-related	AS09 483
AT1G64740	Tubulin alpha chain	AS10 680
AT1G66350	RGL1	AS11 1802
AT1G67090	RbcS Rubisco small subunit (SSU) from pea	AS07 222
AT1G68010	HPR hydroxypyruvate reductase (peroxysomal matrix marker)	AS11 1797
AT1G68830	STN7 serine/threonine-protein kinase STN7, chloroplastic (100 ul)	AS10 1611
AT1G69780	WUSCHEL-related homeobox 4	AS11 1773
AT1G73990	SppA1 signal peptide peptidase (higher plants) (300 ul)	AS10 1598
AT1G74310	HSP101 ClpB heat shock protein, N-terminal	AS07 253
AT1G74310	HSP101 ClpB heat shock protein, N-terminal	AS08 283
AT1G74310	HSP101 ClpB heat shock protein, C-terminal	AS08 287
AT1G74310	ATPase AAA2 domain	AS11 1754
AT1G75270	DHAR2 Dehydroascorbate Reductase 2	AS11 1747
AT1G75820	CLV1 receptor protein kinase CLAVATA1	AS11 1779
AT1G76030	V-ATPase, B vacuolar H ⁺ -ATPase subunit B	AS09 503
AT1G77120	ADH alcohol dehydrogenase (hypoxia marker)	AS10 685
AT1G77490	APX L-ascorbate peroxidase	AS08 368
AT1G77760	NR Nitrate reductase, assimilatory	AS08 310
AT1G78900	V-ATPase, A vacuolar H ⁺ -ATPase subunit A	AS09 467
AT1G78900	V-ATPase, A vacuolar H ⁺ -ATPase subunit A	AS09 502
AT1G79040	PsbR 10 kDa protein of PSII	AS05 059
AT2G05100	Lhcb2 LHCII type II chlorophyll a/b-binding protein	AS01 003
AT2G06520	PsbX small subunit X of PSII	AS05 066
AT2G13540	CBP80 nuclear cap-binding protein subunit 1	AS09 531
AT2G14580	PR-1 Pathogenesis-related protein 1	AS10 687
AT2G16510	V-ATPase, c vacuolar H ⁺ -ATPase, subunit c (16 kDa)	AS09 468
AT2G17950	AtWUS WUSCHEL protein	AS11 1759

TAB]

AT2G18110	eEF1b elongation factor eEF1b-beta protein	AS07 265
AT2G19310	HSP18.5 class IV heat shock protein	AS11 1628
AT2G20890	Psb29 thylakoid membrane formation 1, THF1	AS07 240
AT2G21410	V-ATPase, a vacuolar H ⁺ ATPase, subunit a	AS09 466
AT2G26140	AtFtsH4 plant fts metallo-protease H4	AS07 205
AT2G26690	AJ011604 nitrate transporter	AS09 473
AT2G27040	AGO4 argonaute 4	AS09 617
AT2G27100	Serrate RNA effector molecule	AS09 532
AT2G27880	AGO5 argonaute 5	AS10 671
AT2G28190	CSD2 chloroplastic Cu/Zn superoxide dismutase	AS06 170
AT2G29080	FtsH3 + FtsH10 plant fts metallo-protease H3 + H10	AS07 204
AT2G29410	AtMTP4 metal tolerance protein B	AS09 474
AT2G30390	HemH protoporphyrin ferrochelatase	AS07 228
AT2G30570	PsbW small subunit W of PSII	AS05 060
AT2G30580	E3 ubiquitin protein ligase DRIP2	AS11 1761
AT2G32940	AGO6 argonaute 6	AS10 672
AT2G32950	COP1 constitutive photomorphogenesis protein 1	AS08 342
AT2G35370	GDC-H H protein of glycine decarboxylase complex (GDC)	AS05 074
AT2G36160	S14 40S ribosomal protein S14-1	AS09 477
AT2G36530	Enolase	AS10 651
AT2G36830	AtTIP1	AS09 482
AT2G36830	TIP1;1, TIP1;2 tonoplast intrinsic protein 1-1, 1-2 (gamma)	AS09 493
AT2G36830	AtSIP1	AS09 495
AT2G37170	PIP2;2 plasma membrane aquaporin 2b	AS09 490
AT2G37340	RS-domain	AS11 1805
AT2G37630	AtMYB91 transcription factor AS1	AS11 1765
AT2G39730	RA Rubisco activase	AS10 700
AT2G41220	GOGAT glutamine oxoglutarate aminotransferase	AS07 242
AT2G41560	Ca ²⁺ -ATPase calmodulin-stimulated calcium-ATPase	AS09 486
AT2G46800	AtMTP1 vacuolar Zn ²⁺ /H ⁺ antiporter	AS09 485
AT2G47170	Arf1 ADP-ribosylation factor 1	AS08 325
AT2G47940	Deg2 chloroplastic DegP-type serine protease 2	AS10 1581
AT3G01090	AKIN10 SNF1-related protein kinase catalytic subunit alpha KIN10	AS10 919
AT3G01280	VDAC1 voltage-dependent anion-selective channel protein 1	AS07 212

TAB]

AT3G01480	CYP38 cyclophilin 38, peptidyl-prolyl cis-trans isomerase (100 ul)	AS10 1607
AT3G03250	UGPase UDP-glucose pyrophosphorylase (cytoplasm marker)	AS05 086
AT3G03380	DegP7 Protease Do-like 7	AS11 1756
AT3G03380	Deg9 Protease Do-like 9 (nucleolus marker)	AS11 1785
AT3G03450	RGL2	AS11 1803
AT3G10920	MnSOD manganese superoxide dismutase	AS09 524
AT3G11130	Clathrin heavy-chain	AS10 690
AT3G12110	Actin-11	AS10 702
AT3G12280	RBR1 retinoblastoma related protein	AS11 1627
AT3G12580	HSP70 heat shock protein 70 cytoplasmic	AS08 371
AT3G15520	PPIase CYP37 peptidyl-prolyl cis-trans isomerase CYP37, chloroplastic (100 ul)	AS10 1589
AT3G16140	PsaH PSI-H subunit of photosystem I	AS06 105
AT3G19900	CPLD11	AS10 937
AT3G20740	Polycomb group protein FERTILIZATION-INDEPENDENT ENDOSPERM	AS11 1760
AT3G21720	ICL isocitrate lyase	AS09 500
AT3G22370	AOX1/2 plant alternative oxidase 1 and 2	AS04 054
AT3G22840	Elip1 early light inducible protein 1	AS03 033
AT3G23890	TOP2 DNA topoisomerase II	AS05 096
AT3G24430	Hcf101 high chlorophyll fluorescence phenotype protein	AS06 163
AT3G25940	A12.2 RNA polymerase I subunit (homolog of Pol II Rpb9)	AS07 225
AT3G26060	PrxQ peroxiredoxin, thioredoxin reductase	AS05 093
AT3G26520	TIP1;1,TIP1;2 tonoplast intrinsic protein1-1, 1-2 (gamma)	AS09 492
AT3G27925	Deg1 chloroplastic DegP-type serine protease 1	AS07 246
AT3G42050	V-ATPase, H vacuolar H ⁺ -ATPase subunit H	AS09 499
AT3G44750	HDT1 histone deacetylase	AS11 1792
AT3G44880	Acd1 accelerated cell death 1	AS11 1783
AT3G45140	LOX-C chloroplastic lipoxygenase	AS07 258
AT3G45780	PHOT1 phototropin-1	AS10 720
AT3G46740	Toc75 chloroplast outer envelope membrane translocon complex OEP75 protein	AS06 150
AT3G46740	Toc75 Protein TOC75-3, chloroplastic, POTRA domain 3	AS08 351
AT3G47470	Lhca4 PSI type IV chlorophyll a/b-binding protein	AS01 008
AT3G48530	AKING1 SNF1-related protein kinase regulatory subunit gamma 1	AS09 613
AT3G48750	CDC2 cell-division-cycle kinase 2	AS06 153
AT3G49010	L13-1 60S ribosomal protein L13-1	AS09 478

TAB]

AT3G50980	Dehydrin	AS07 206
AT3G52750	FtsZ plant cell division protein ftsZ	AS09 413
AT3G52930	ALD fructose-1,6 bisphosphate aldolase	AS08 294
AT3G53420	PIP2;1 aquaporin PIP2;1	AS09 488
AT3G53420	PIP2;1, PIP2;2, PIP2;3 plasma membrane intrinsic protein 2-1,2-2,2-3	AS09 491
AT3G54030	Protein kinase protein with tetratricopeptide repeat domain	AS11 1793
AT3G54660	GR glutathione reductase	AS06 181
AT3G54890	Lhca1 PSI type I chlorophyll a/b-binding protein	AS01 005
AT3G56940	CRD1 cyanobacterial homolog of plant CHL27 cyclase	AS06 122
AT3G56950	AtSIP2	AS09 496
AT3G58730	V-ATPase, D vacuolar H ⁺ -ATPase subunit D	AS09 497
AT3G61430	PIP1 (PIP1;1, PIP1;2, PIP1;3) aquaporins	AS09 487
AT3G61430	PIP1;1, PIP1;2, PIP1;3 aquaporins	AS09 489
AT3G61470	Lhca2 PSI type II chlorophyll a/b-binding protein	AS01 006
AT3G62560	Sar1 secretion-associated and Ras-related protein 1	AS08 326
AT4G02390	PARP-2 Poly [ADP-ribose] polymerase 2	AS10 675
AT4G02520	GST class-phi glutathione S tranferase	AS09 479
AT4G02630	TAK1 serine/threonine protein kinase (300 ul)	AS10 1584
AT4G02630	TAK1 serine/threonine protein kinase (100 ul)	AS10 1612
AT4G03280	PetC Rieske iron-sulfur protein of Cyt b6/f complex	AS08 330
AT4G04020	PGL35 plastoglobulin 35	AS06 116
AT4G04640	AtpC gamma subunit of ATP synthase	AS08 312
AT4G05050	UBQ11 Ubiquitin	AS08 307
AT4G08150	homeodomain protein KNAT1/BP	AS11 1763
AT4G09650	AtpD CF1 delta subunit of ATP synthase, Serum	AS10 1591
AT4G10340	Lhcb5 CP26 chlorophyll a/b-binding protein of plant PSII	AS01 009
AT4G11010	NDPK nucleoside diphosphate kinase	AS08 302
AT4G11150	V-ATPase epsilon subunit of tonoplast H ⁺ ATPase (200 µl)	AS07 213
AT4G11150	V-ATPase epsilon subunit of tonoplast H ⁺ ATPase	AS09 577
AT4G12800	PsaL PSI-L subunit of photosystem I	AS06 108
AT4G14690	Elip2 early light inducible protein 2	AS03 036
AT4G16360	AKINB2 SNF1-related protein kinase regulatory subunit beta-2	AS09 462
AT4G17340	TIP2;2 tonoplast intrinsic protein 2-2	AS09 510
AT4G18370	Deg5 protease Do-like 5, chloroplastic	AS10 703

TAB]

AT4G19690	IRT1 iron regulated transporter 1	AS11 1780
AT4G20260	AtPCaP1 Arabidopsis thaliana plasma membrane cation-binding protein-1	AS10 931
AT4G21280	PsbQ 16 kDa protein of the oxygen evolving complex (OEC) of PSII	AS06 142-16
AT4G22890	PGRL1 PGR5-like protein 1A, chloroplastic	AS10 725
AT4G23100	Gamma-ECS gamma glutamylcysteine synthase	AS06 186
AT4G24280	HSP70 heat shock protein 70, chloroplastic	AS08 348
AT4G25100	FeSOD chloroplastic Fe-dependent superoxide dismutase	AS06 125
AT4G26840	SUMO1 small ubiquitin-like modifier protein 1	AS08 308
AT4G26850	GDP-L-Galactose Phosphorylase	AS10 723
AT4G27650	PEL1 Pelota	AS11 1798
AT4G27670	HSP21 chloroplastic heat shock protein	AS08 285
AT4G28750	PsaE PSI-E subunit of photosystem I	AS08 324
AT4G31870	GPX chloroplastic glutathione peroxidase	AS04 055
AT4G32650	KC1 potassium channel KAT3	AS09 515
AT4G34450	Sec21p gamma subunit, COP vesicles	AS08 327
AT4G35090	Catalase	AS09 501
AT4G35100	PIP2-7 plasma membrane aquaporin	AS09 469
AT4G35260	Idh isocitrate dehydrogenase	AS06 203A
AT4G35800	RNA polymerase II subunit B1	AS11 1804
AT4G35830	ACO1 aconitase	AS09 521
AT4G37910	HSP90-1 heat shock protein 90-1	AS08 346
AT4G37910	HSP70 heat shock protein 70, mitochondrial	AS08 347
AT4G37930	SHMT serine hydroxymethyltransferase	AS05 075
AT4G38630	DS5a Drosophila 26S proteasome subunit Rpn10	AS01 012
AT4G40030	H3 histone H3	AS10 710
AT5G01530	Lhcb4 CP29 chlorophyll a/b binding protein of plant PSII	AS04 045
AT5G01600	Ferritin (plant)	AS10 674
AT5G01920	STN8 kinase serine/threonine-protein kinase STN8, chloroplastic (100 ul)	AS10 1601
AT5G03740	HDT3 histone deacetylase	AS11 1753
AT5G05660	NFX2 nuclear transcription factor, X-box-binding protein	AS10 701
AT5G05660	NFX2 nuclear transcription factor, x-box binding protein	AS11 1755
AT5G12110	eEF1B-alpha1 an 2 elongation factor 1B-alpha 1 and 1B-alpha 2	AS10 678
AT5G13710	SMT1 Sterol methyltransferase 1	AS07 266
AT5G15250	FtsH6 plant fts metallo-protease H6	AS05 094A

TAB]

AT5G15450	ClpB-P ClpB3	AS09 459
AT5G16620	Tic40 chloroplast inner envelope membrane translocon complex protein (200 µl)	AS10 709
AT5G18170	GDH1 glutamate dehydrogenase 1	AS05 068
AT5G19510	eEF1B-alpha2 elongation factor 1B-alpha 2	AS10 679
AT5G20280	SPS sucrose phosphate synthase, global	AS03 035A
AT5G21150	AGO9 argonaute 9	AS10 673
AT5G21170	AKINB1 SNF1-related protein kinase regulatory subunit beta-1	AS09 460
AT5G27150	NHE-1 Na ⁺ /H ⁺ antiporter, sodium/hydrogen exchanger	AS09 484
AT5G27380	GS glutathione synthase, GSH-S	AS06 183
AT5G35630	GLN2 GS2, chloroplastic form of glutamine synthetase	AS08 296
AT5G37600	GLN1 GLN2 GS1 glutamine synthetase global antibody	AS08 295
AT5G38430	RbcS Rubisco small subunit (SSU)	AS07 259
AT5G42020	BiP2 lumenal-binding protein 2	AS09 481
AT5G43940	GSNOR S-nitrosoglutathione reductase	AS09 647
AT5G44200	CBP20 nuclear cap-binding protein subunit 2	AS09 530
AT5G48300	ADGP ADP-glucose pyrophosphorylase	AS11 1739
AT5G49470	Hik29 sensor histidine kinase (300 ul)	AS10 1599
AT5G49470	Hik29 sensor histidine kinase (100 ul)	AS10 1600
AT5G50920	ClpC chloroplastic form of HSP100	AS01 001
AT5G54270	Lhcb3 LHCII type III chlorophyll a/b-binding protein	AS01 002
AT5G54960	PDC pyruvate decarboxylase	AS10 691
AT5G56030	HSP90-2 heat shock protein 90-2	AS11 1629
AT5G57090	PIN2 auxin efflux carrier PIN2	AS05 069
AT5G58140	PHOT2 phototropin-2	AS10 721
AT5G58250	CGL78 Genomic DNA, chromosome 5, P1 clone:MCK7	AS10 936
AT5G58870	FtsH9 plant fts metallo-protease H9	AS06 130
AT5G62690	Tubulin beta chain	AS10 681
AT5G62810	AtPex14p peroxysomal marker	AS08 372
AT5G63980	SAL1 sal1 phosphatase	AS07 256
AT5G64040	PsaN PSI-N subunit of photosystem I	AS06 109
AT5G66570	PsbO 33 kDa of the oxygen evolving complex (OEC) of PSII	AS05 092
AT5G67030	ZEP Zeaxanthin Epoxidase	AS08 289
ATCG00020	PsbA D1 protein of PSII, C-terminal (10 µl)	AS05 084-10
ATCG00080	PsbI small subunit I of PSII	AS06 158

TAB]

ATCG00120	AtpA alpha subunit of ATP synthase	AS08 304
ATCG00140	AtpH ATP synthase subunit c	AS05 071
ATCG00140	AtpH ATP synthase subunit c	AS09 591
ATCG00270	PsbD D2 protein of PSII	AS06 146
ATCG00280	PsbC CP43 protein of PSII	AS06 110
ATCG00300	PsbZ ycf9 protein of PSII	AS06 115
ATCG00340	PsaB PSI-B core subunit of photosystem I	AS10 695
ATCG00350	PsaA PSI-A core protein of photosystem I	AS06 172
ATCG00350	cAPX cytosolic ascorbate peroxidase	AS06 180
ATCG00490	RbcL Rubisco large subunit, form I	AS01 017
ATCG00490	RbcL Rubisco large subunit, form I and form II (100 µg)	AS03 037
ATCG00540	Cyt f cytochrome f protein (PetA) of thylakoid Cyt b6/f-complex	AS06 119
ATCG00570	PsbF beta subunit of Cytochrome b559 of PSII	AS06 113
ATCG00580	PsbE alfa subunit of Cytochrome b559 of PSII	AS06 112
ATCG00680	PsbB CP47 protein of PSII	AS04 038
ATCG00710	PsbH small subunit H of PSII	AS06 157
ATCG00720	Cyt b6 thylakoid membrane cytochrome b6 protein, N terminal	AS03 034
ATCG01060	PsaC PSI-C core subunit of photosystem I	AS10 939
ATMG00160	COXII cytochrome oxidase subunit II	AS04 053A

TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

URL to product page

<http://www.agrisera.com/AT1G02280>
<http://www.agrisera.com/AT1G03130>
<http://www.agrisera.com/AT1G06680>
<http://www.agrisera.com/AT1G06760.1>
<http://www.agrisera.com/AT1G07510>
<http://www.agrisera.com/AT1G07940>
<http://www.agrisera.com/AT1G08380>
<http://www.agrisera.com/AT1G09340>
<http://www.agrisera.com/AT1G09570>
<http://www.agrisera.com/AT1G09640>
<http://www.agrisera.com/AT1G09700>
<http://www.agrisera.com/AT1G09940>
<http://www.agrisera.com/AT1G11510>
<http://www.agrisera.com/AT1G12030>
<http://www.agrisera.com/AT1G12520>
<http://www.agrisera.com/AT1G15690>
<http://www.agrisera.com/AT1G15820>
<http://www.agrisera.com/AT1G19570>
<http://www.agrisera.com/AT1G20340>
<http://www.agrisera.com/AT1G22840>
<http://www.agrisera.com/AT1G23490>
<http://www.agrisera.com/AT1G28320>
<http://www.agrisera.com/AT1G29910>
<http://www.agrisera.com/AT1G30230>
<http://www.agrisera.com/AT1G30380>
<http://www.agrisera.com/AT1G30400>
<http://www.agrisera.com/AT1G31330>
<http://www.agrisera.com/AT1G31810>
<http://www.agrisera.com/AT1G32060>
<http://www.agrisera.com/AT1G43670>
<http://www.agrisera.com/AT1G44575>
<http://www.agrisera.com/AT1G45474>
<http://www.agrisera.com/AT1G48410>

TAB]

<http://www.agrisera.com/AT1G53540>
<http://www.agrisera.com/AT1G54780>
<http://www.agrisera.com/AT1G55670>
<http://www.agrisera.com/AT1G59870>
<http://www.agrisera.com/AT1G61520>
<http://www.agrisera.com/AT1G62300>
<http://www.agrisera.com/AT1G62360>
<http://www.agrisera.com/AT1G62480>
<http://www.agrisera.com/AT1G64740>
<http://www.agrisera.com/AT1G66350>
<http://www.agrisera.com/AT1G67090>
<http://www.agrisera.com/AT1G68010>
<http://www.agrisera.com/AT1G68830>
<http://www.agrisera.com/AT1G69780>
<http://www.agrisera.com/AT1G73990>
<http://www.agrisera.com/AT1G74310>
<http://www.agrisera.com/AT1G74310>
<http://www.agrisera.com/AT1G74310>
<http://www.agrisera.com/AT1G74310>
<http://www.agrisera.com/AT1G75270>
<http://www.agrisera.com/AT1G75820>
<http://www.agrisera.com/AT1G76030>
<http://www.agrisera.com/AT1G77120>
<http://www.agrisera.com/AT1G77490>
<http://www.agrisera.com/AT1G77760>
<http://www.agrisera.com/AT1G78900>
<http://www.agrisera.com/AT1G78900>
<http://www.agrisera.com/AT1G79040>
<http://www.agrisera.com/AT2G05100>
<http://www.agrisera.com/AT2G06520>
<http://www.agrisera.com/AT2G13540>
<http://www.agrisera.com/AT2G14580>
<http://www.agrisera.com/AT2G16510>
<http://www.agrisera.com/AT2G17950>

TAB]

<http://www.agrisera.com/AT2G18110>
<http://www.agrisera.com/AT2G19310>
<http://www.agrisera.com/AT2G20890>
<http://www.agrisera.com/AT2G21410>
<http://www.agrisera.com/AT2G26140>
<http://www.agrisera.com/AT2G26690>
<http://www.agrisera.com/AT2G27040>
<http://www.agrisera.com/AT2G27100>
<http://www.agrisera.com/AT2G27880>
<http://www.agrisera.com/AT2G28190>
<http://www.agrisera.com/AT2G29080>
<http://www.agrisera.com/AT2G29410>
<http://www.agrisera.com/AT2G30390>
<http://www.agrisera.com/AT2G30570>
<http://www.agrisera.com/AT2G30580>
<http://www.agrisera.com/AT2G32940>
<http://www.agrisera.com/AT2G32950>
<http://www.agrisera.com/AT2G35370>
<http://www.agrisera.com/AT2G36160>
<http://www.agrisera.com/AT2G36530>
<http://www.agrisera.com/AT2G36830>
<http://www.agrisera.com/AT2G36830>
<http://www.agrisera.com/AT2G36830>
<http://www.agrisera.com/AT2G37170>
<http://www.agrisera.com/AT2G37340>
<http://www.agrisera.com/AT2G37630>
<http://www.agrisera.com/AT2G39730>
<http://www.agrisera.com/AT2G41220>
<http://www.agrisera.com/AT2G41560>
<http://www.agrisera.com/AT2G46800>
<http://www.agrisera.com/AT2G47170>
<http://www.agrisera.com/AT2G47940>
<http://www.agrisera.com/AT3G01090>
<http://www.agrisera.com/AT3G01280>

TAB]

<http://www.agrisera.com/AT3G01480>
<http://www.agrisera.com/AT3G03250>
<http://www.agrisera.com/AT3G03380>
<http://www.agrisera.com/AT3G03380>
<http://www.agrisera.com/AT3G03450>
<http://www.agrisera.com/AT3G10920>
<http://www.agrisera.com/AT3G11130>
<http://www.agrisera.com/AT3G12110>
<http://www.agrisera.com/AT3G12280>
<http://www.agrisera.com/AT3G12580>
<http://www.agrisera.com/AT3G15520>
<http://www.agrisera.com/AT3G16140>
<http://www.agrisera.com/AT3G19900>
<http://www.agrisera.com/AT3G20740>
<http://www.agrisera.com/AT3G21720>
<http://www.agrisera.com/AT3G22370>
<http://www.agrisera.com/AT3G22840>
<http://www.agrisera.com/AT3G23890>
<http://www.agrisera.com/AT3G24430>
<http://www.agrisera.com/AT3G25940>
<http://www.agrisera.com/AT3G26060>
<http://www.agrisera.com/AT3G26520>
<http://www.agrisera.com/AT3G27925>
<http://www.agrisera.com/AT3G42050>
<http://www.agrisera.com/AT3G44750>
<http://www.agrisera.com/AT3G44880>
<http://www.agrisera.com/AT3G45140>
<http://www.agrisera.com/AT3G45780>
<http://www.agrisera.com/AT3G46740>
<http://www.agrisera.com/AT3G46740>
<http://www.agrisera.com/AT3G47470>
<http://www.agrisera.com/AT3G48530>
<http://www.agrisera.com/AT3G48750>
<http://www.agrisera.com/AT3G49010>

TAB]

<http://www.agrisera.com/AT3G50980>
<http://www.agrisera.com/AT3G52750>
<http://www.agrisera.com/AT3G52930>
<http://www.agrisera.com/AT3G53420>
<http://www.agrisera.com/AT3G53420>
<http://www.agrisera.com/AT3G54030>
<http://www.agrisera.com/AT3G54660>
<http://www.agrisera.com/AT3G54890>
<http://www.agrisera.com/AT3G56940>
<http://www.agrisera.com/AT3G56950>
<http://www.agrisera.com/AT3G58730>
<http://www.agrisera.com/AT3G61430>
<http://www.agrisera.com/AT3G61430>
<http://www.agrisera.com/AT3G61470>
<http://www.agrisera.com/AT3G62560>
<http://www.agrisera.com/AT4G02390>
<http://www.agrisera.com/AT4G02520>
<http://www.agrisera.com/AT4G02630>
<http://www.agrisera.com/AT4G02630>
<http://www.agrisera.com/AT4G03280>
<http://www.agrisera.com/AT4G04020>
<http://www.agrisera.com/AT4G04640>
<http://www.agrisera.com/AT4G05050>
<http://www.agrisera.com/AT4G08150>
<http://www.agrisera.com/AT4G09650>
<http://www.agrisera.com/AT4G10340>
<http://www.agrisera.com/AT4G11010>
<http://www.agrisera.com/AT4G11150>
<http://www.agrisera.com/AT4G11150>
<http://www.agrisera.com/AT4G12800>
<http://www.agrisera.com/AT4G14690>
<http://www.agrisera.com/AT4G16360>
<http://www.agrisera.com/AT4G17340>
<http://www.agrisera.com/AT4G18370>

TAB]

<http://www.agrisera.com/AT4G19690>
<http://www.agrisera.com/AT4G20260>
<http://www.agrisera.com/AT4G21280>
<http://www.agrisera.com/AT4G22890>
<http://www.agrisera.com/AT4G23100>
<http://www.agrisera.com/AT4G24280>
<http://www.agrisera.com/AT4G25100>
<http://www.agrisera.com/AT4G26840>
<http://www.agrisera.com/AT4G26850>
<http://www.agrisera.com/AT4G27650>
<http://www.agrisera.com/AT4G27670>
<http://www.agrisera.com/AT4G28750>
<http://www.agrisera.com/AT4G31870>
<http://www.agrisera.com/AT4G32650>
<http://www.agrisera.com/AT4G34450>
<http://www.agrisera.com/AT4G35090>
<http://www.agrisera.com/AT4G35100>
<http://www.agrisera.com/AT4G35260>
<http://www.agrisera.com/AT4G35800>
<http://www.agrisera.com/AT4G35830>
<http://www.agrisera.com/AT4G37910>
<http://www.agrisera.com/AT4G37910>
<http://www.agrisera.com/AT4G37930>
<http://www.agrisera.com/AT4G38630>
<http://www.agrisera.com/AT4G40030>
<http://www.agrisera.com/AT5G01530>
<http://www.agrisera.com/AT5G01600>
<http://www.agrisera.com/AT5G01920>
<http://www.agrisera.com/AT5G03740>
<http://www.agrisera.com/AT5G05660>
<http://www.agrisera.com/AT5G05660>
<http://www.agrisera.com/AT5G12110>
<http://www.agrisera.com/AT5G13710>
<http://www.agrisera.com/AT5G15250>

TAB]

<http://www.agrisera.com/AT5G15450>
<http://www.agrisera.com/AT5G16620>
<http://www.agrisera.com/AT5G18170>
<http://www.agrisera.com/AT5G19510>
<http://www.agrisera.com/AT5G20280>
<http://www.agrisera.com/AT5G21150>
<http://www.agrisera.com/AT5G21170>
<http://www.agrisera.com/AT5G27150>
<http://www.agrisera.com/AT5G27380>
<http://www.agrisera.com/AT5G35630>
<http://www.agrisera.com/AT5G37600>
<http://www.agrisera.com/AT5G38430>
<http://www.agrisera.com/AT5G42020>
<http://www.agrisera.com/AT5G43940>
<http://www.agrisera.com/AT5G44200>
<http://www.agrisera.com/AT5G48300>
<http://www.agrisera.com/AT5G49470>
<http://www.agrisera.com/AT5G49470>
<http://www.agrisera.com/AT5G50920>
<http://www.agrisera.com/AT5G54270>
<http://www.agrisera.com/AT5G54960>
<http://www.agrisera.com/AT5G56030>
<http://www.agrisera.com/AT5G57090>
<http://www.agrisera.com/AT5G58140>
<http://www.agrisera.com/AT5G58250>
<http://www.agrisera.com/AT5G58870>
<http://www.agrisera.com/AT5G62690>
<http://www.agrisera.com/AT5G62810>
<http://www.agrisera.com/AT5G63980>
<http://www.agrisera.com/AT5G64040>
<http://www.agrisera.com/AT5G66570>
<http://www.agrisera.com/AT5G67030>
<http://www.agrisera.com/ATCG00020>
<http://www.agrisera.com/ATCG00080>

TAB]

<http://www.agrisera.com/ATCG00120>
<http://www.agrisera.com/ATCG00140>
<http://www.agrisera.com/ATCG00140>
<http://www.agrisera.com/ATCG00270>
<http://www.agrisera.com/ATCG00280>
<http://www.agrisera.com/ATCG00300>
<http://www.agrisera.com/ATCG00340>
<http://www.agrisera.com/ATCG00350>
<http://www.agrisera.com/ATCG00350>
<http://www.agrisera.com/ATCG00490>
<http://www.agrisera.com/ATCG00490>
<http://www.agrisera.com/ATCG00540>
<http://www.agrisera.com/ATCG00570>
<http://www.agrisera.com/ATCG00580>
<http://www.agrisera.com/ATCG00680>
<http://www.agrisera.com/ATCG00710>
<http://www.agrisera.com/ATCG00720>
<http://www.agrisera.com/ATCG01060>
<http://www.agrisera.com/ATMG00160>

TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

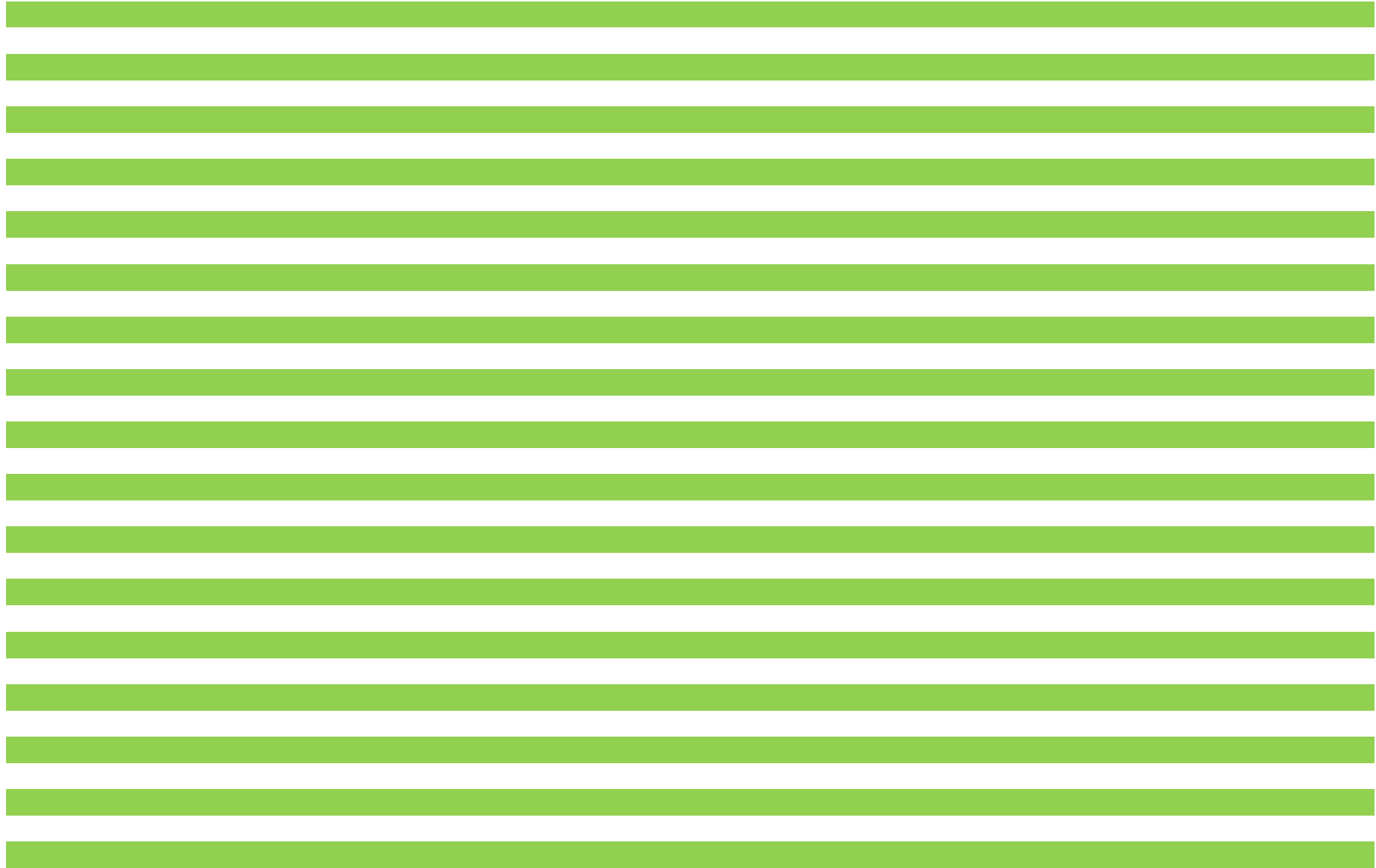
TAB]

TAB]

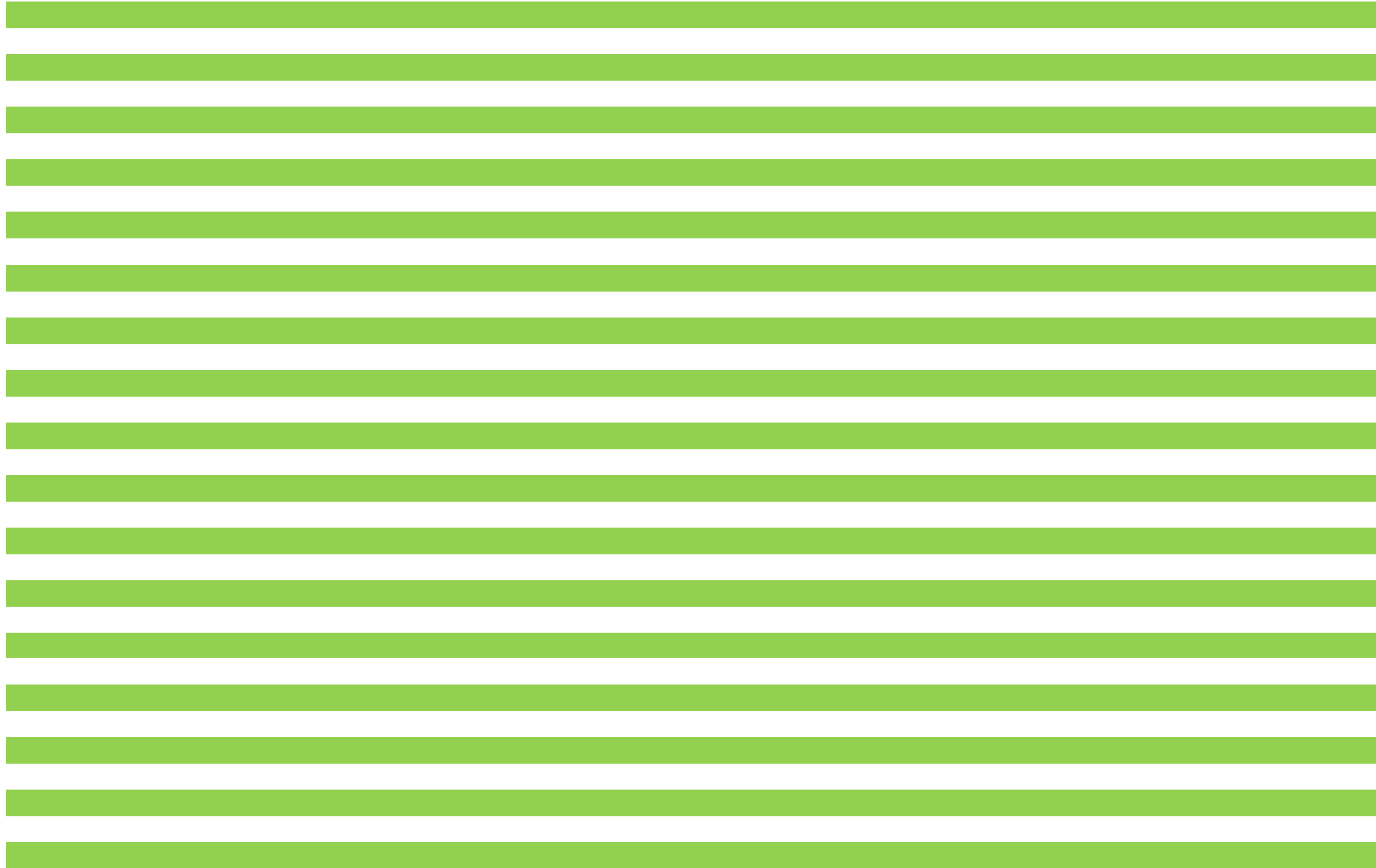
TAB]

TAB]

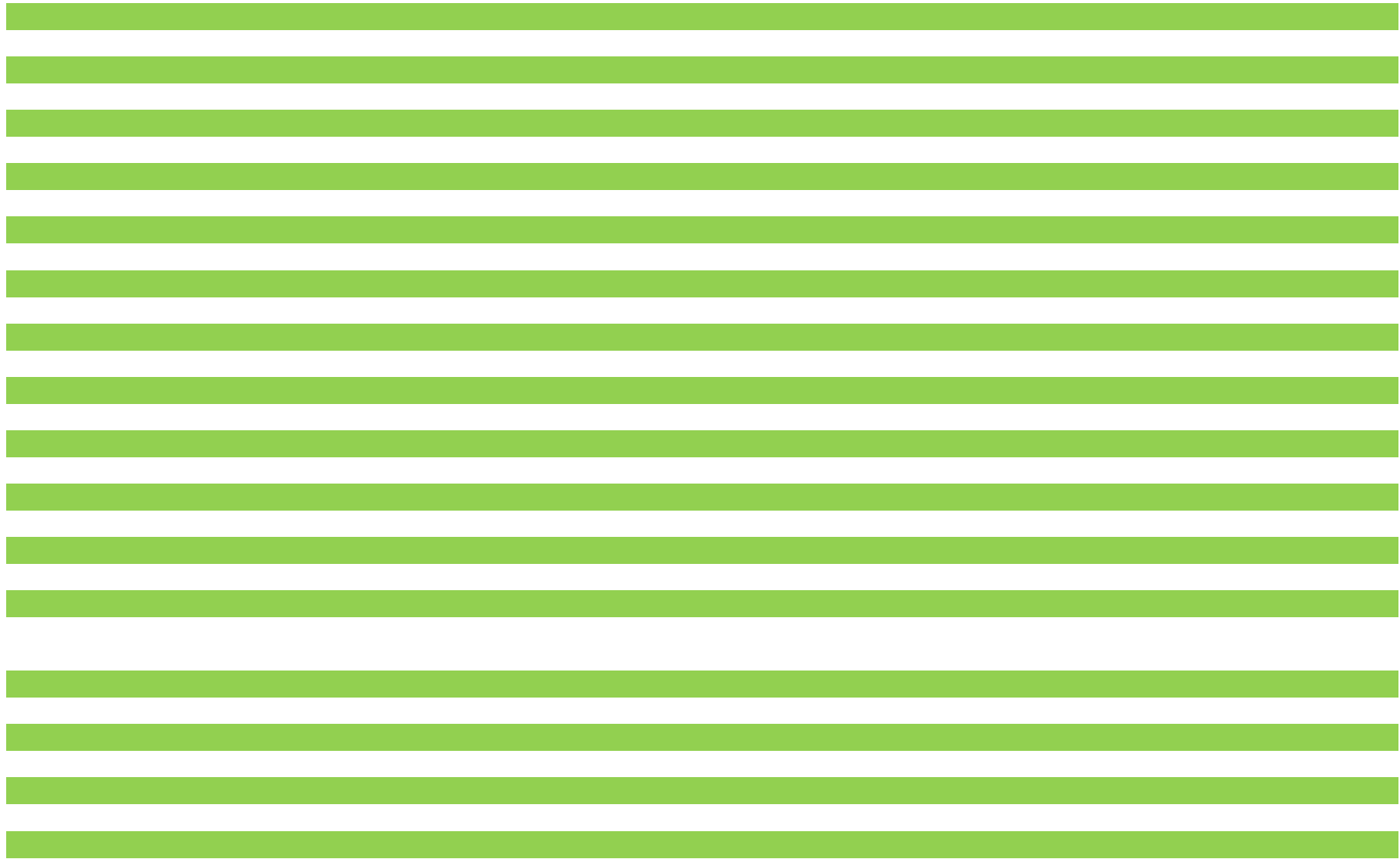
TAB]



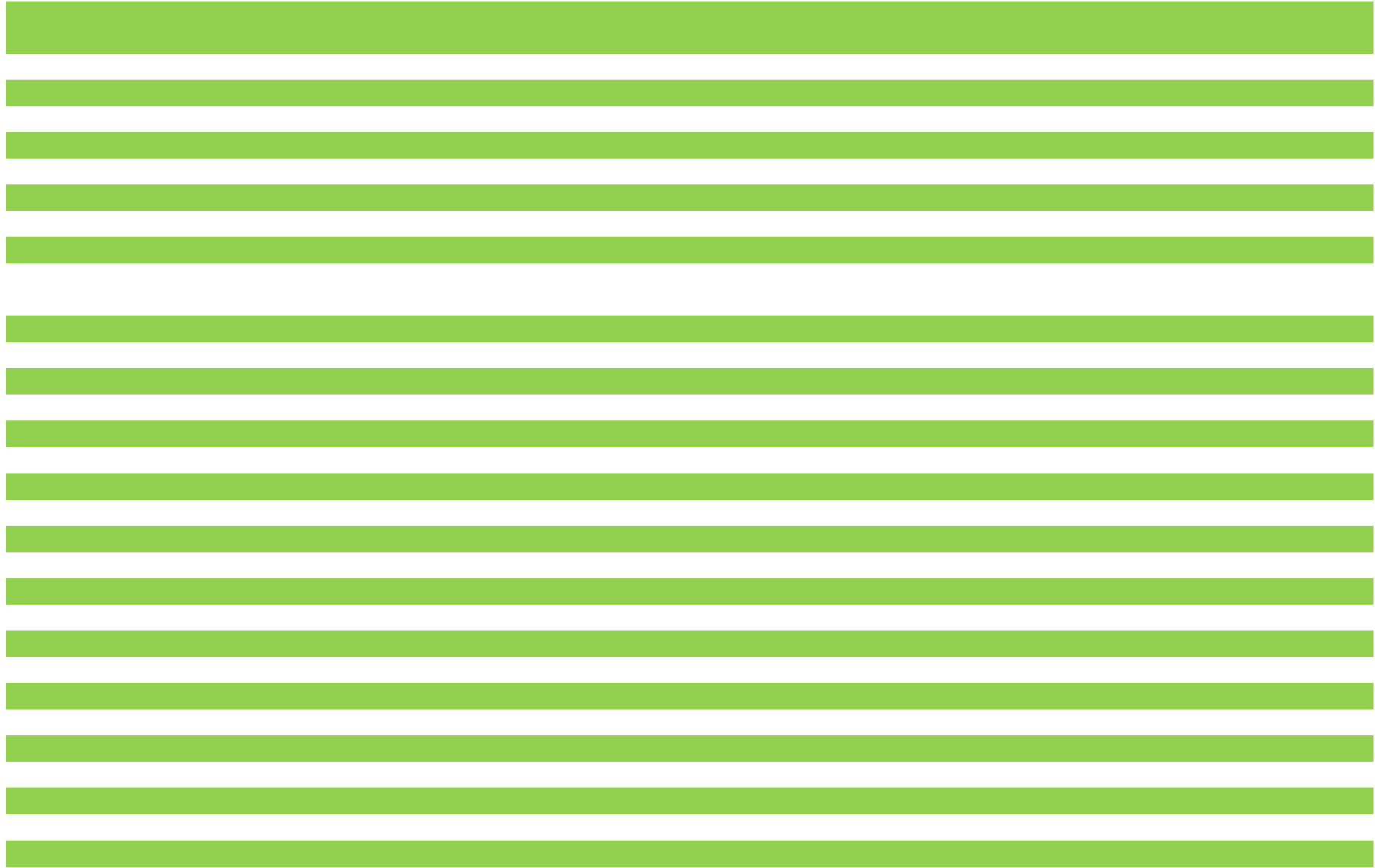
TAB]



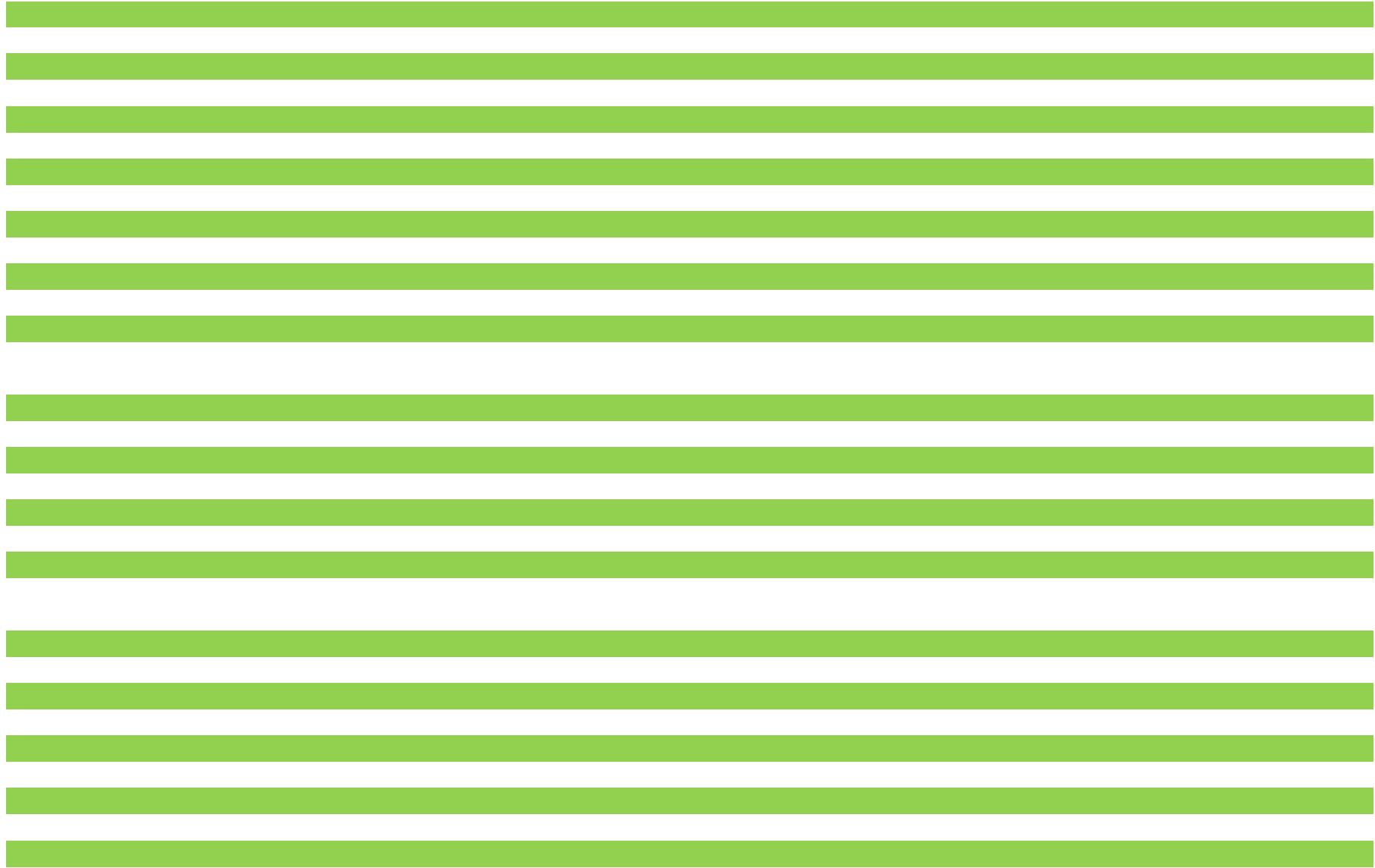
TAB]



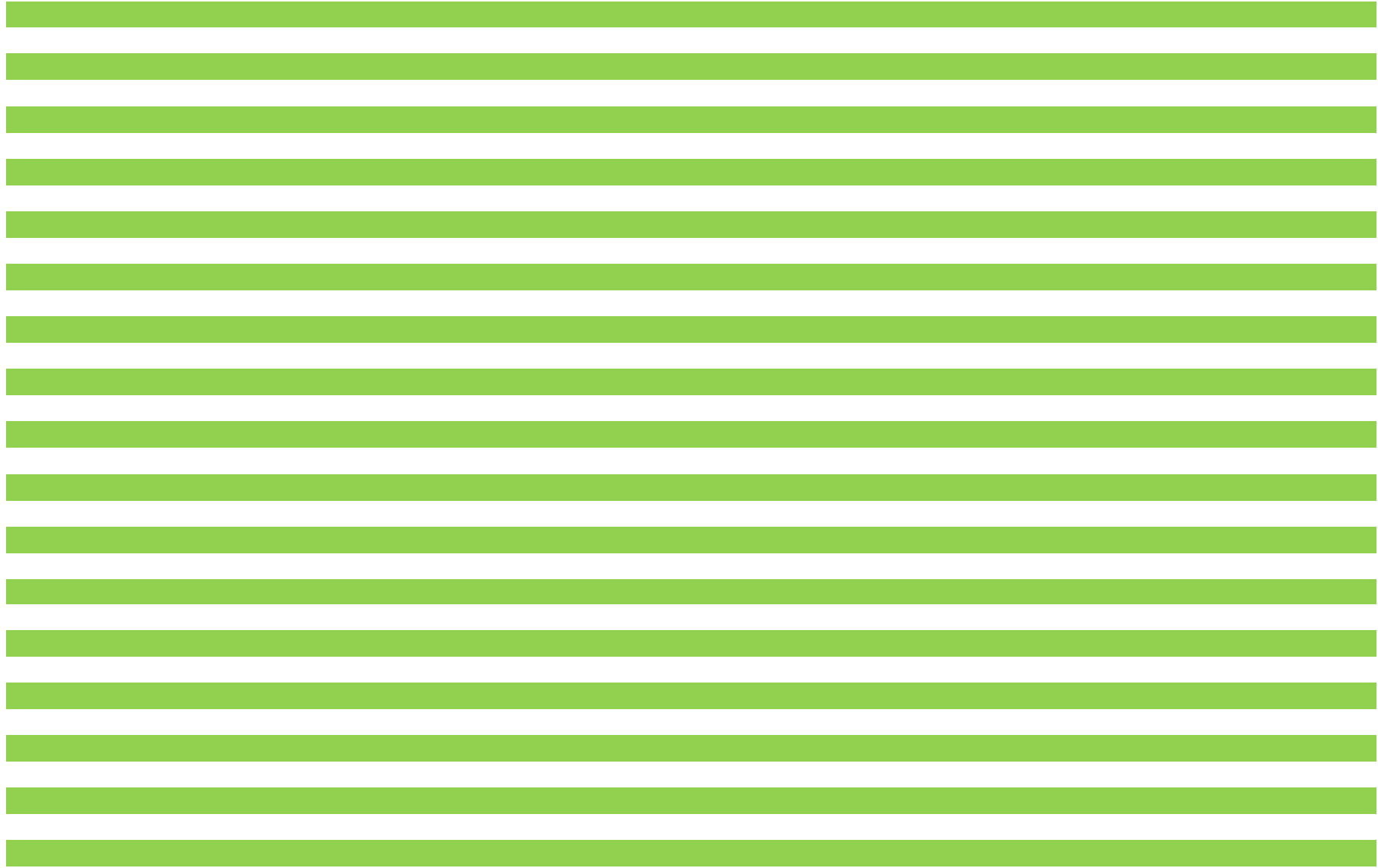
TAB]



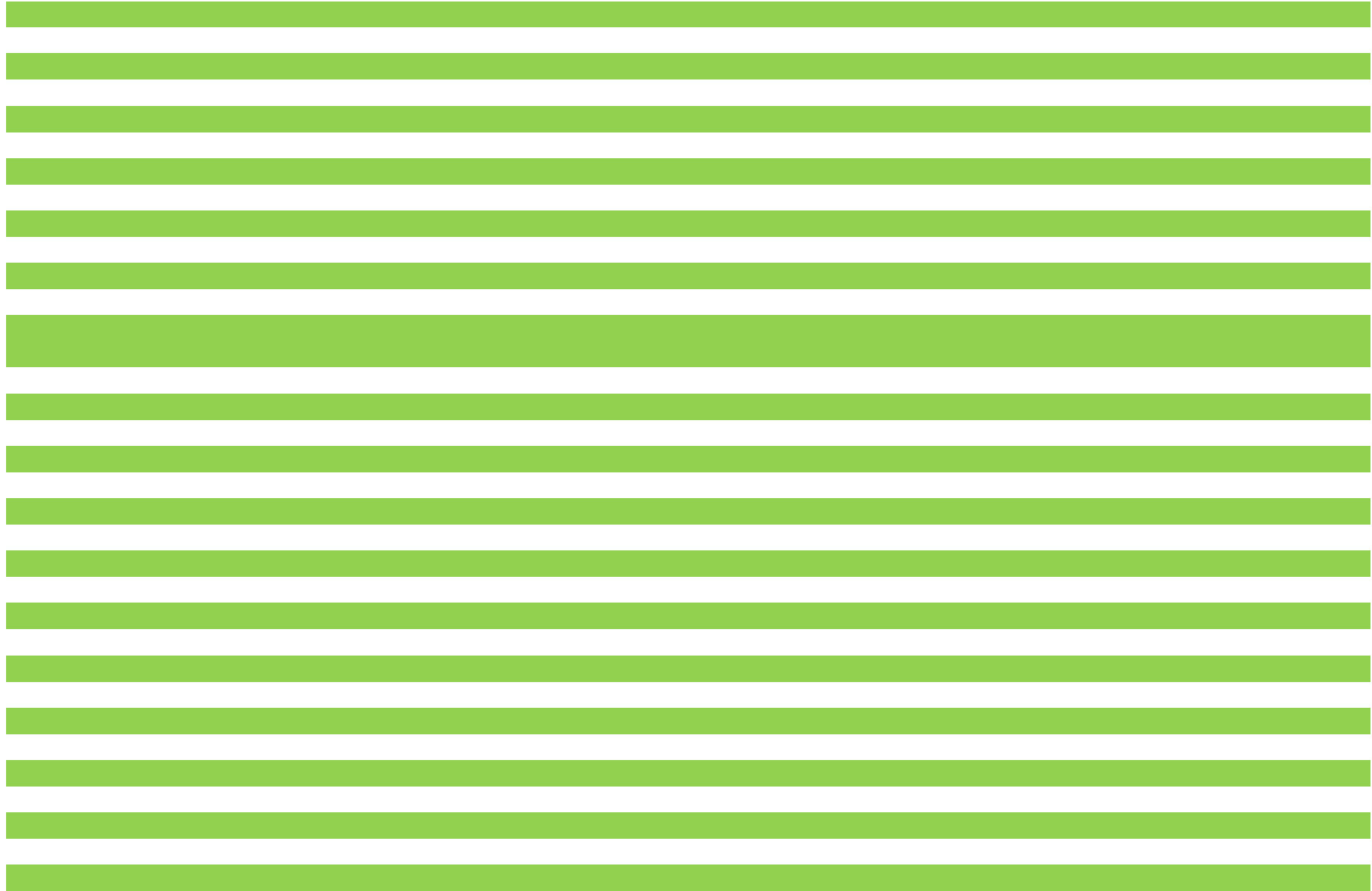
TAB]



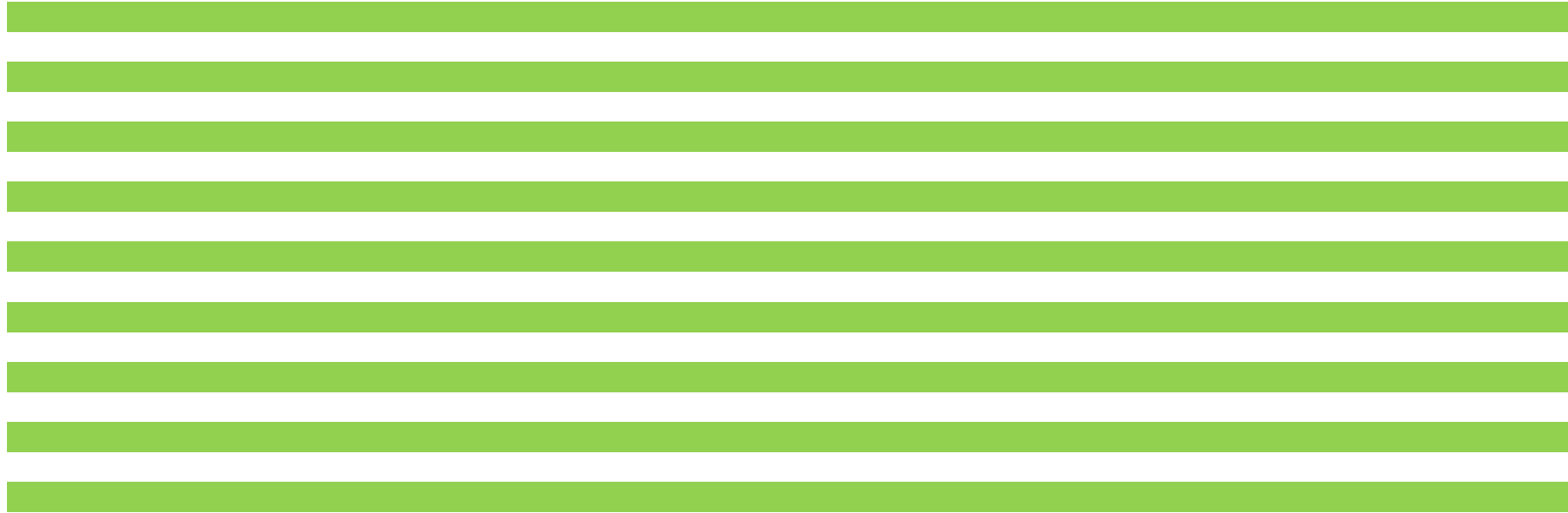
TAB]



TAB]



TAB]



TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

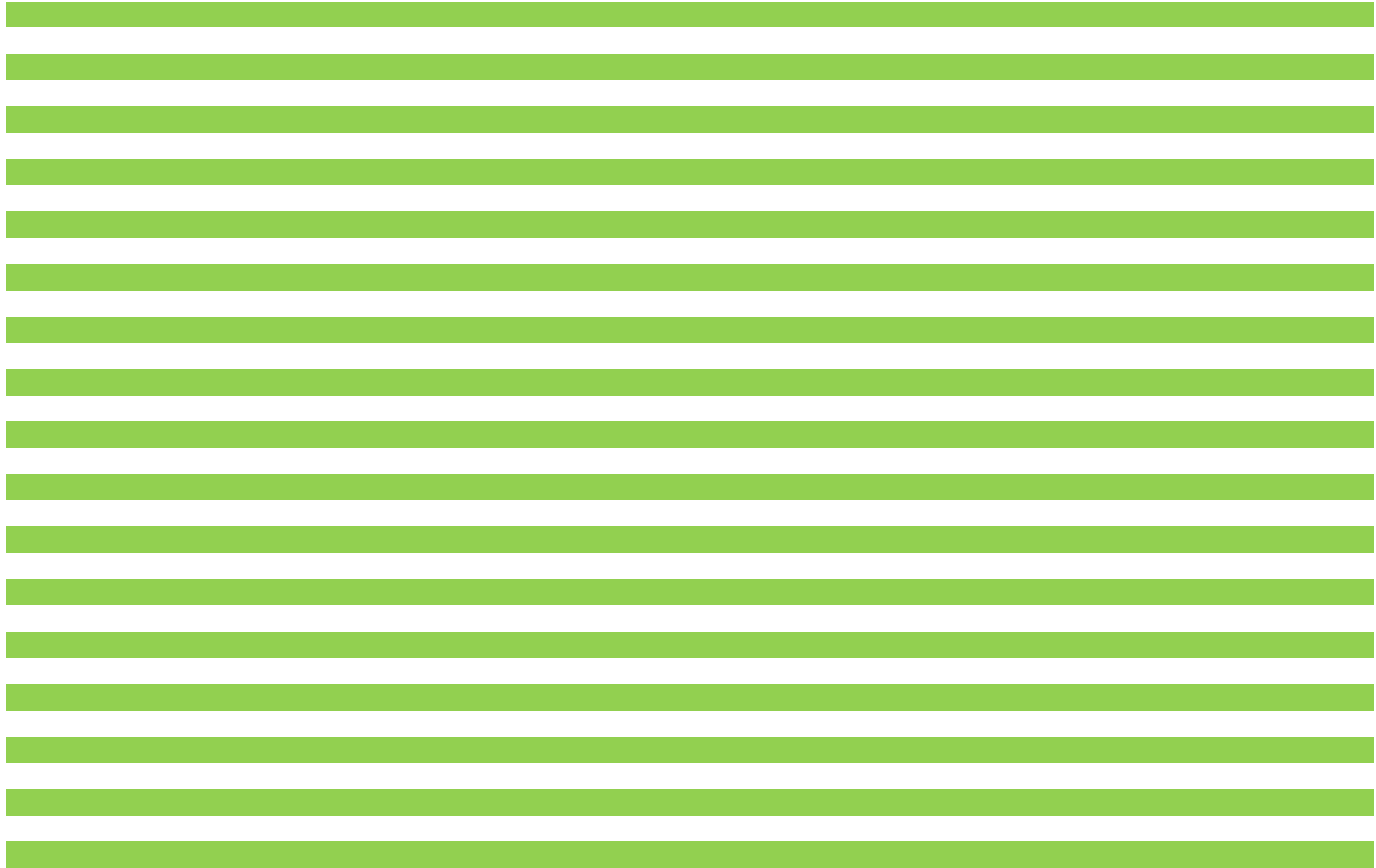
TAB]

TAB]

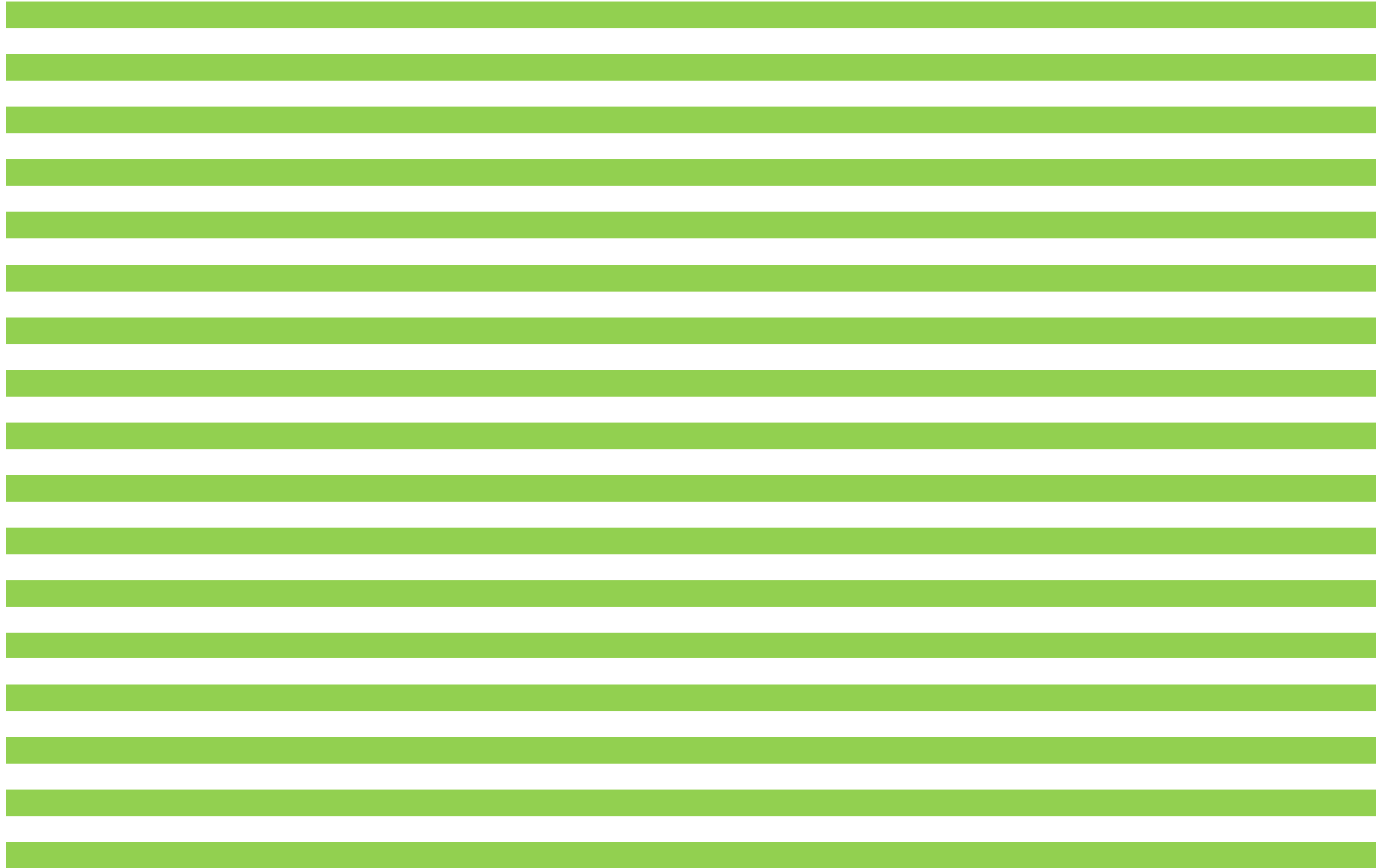
TAB]

TAB]

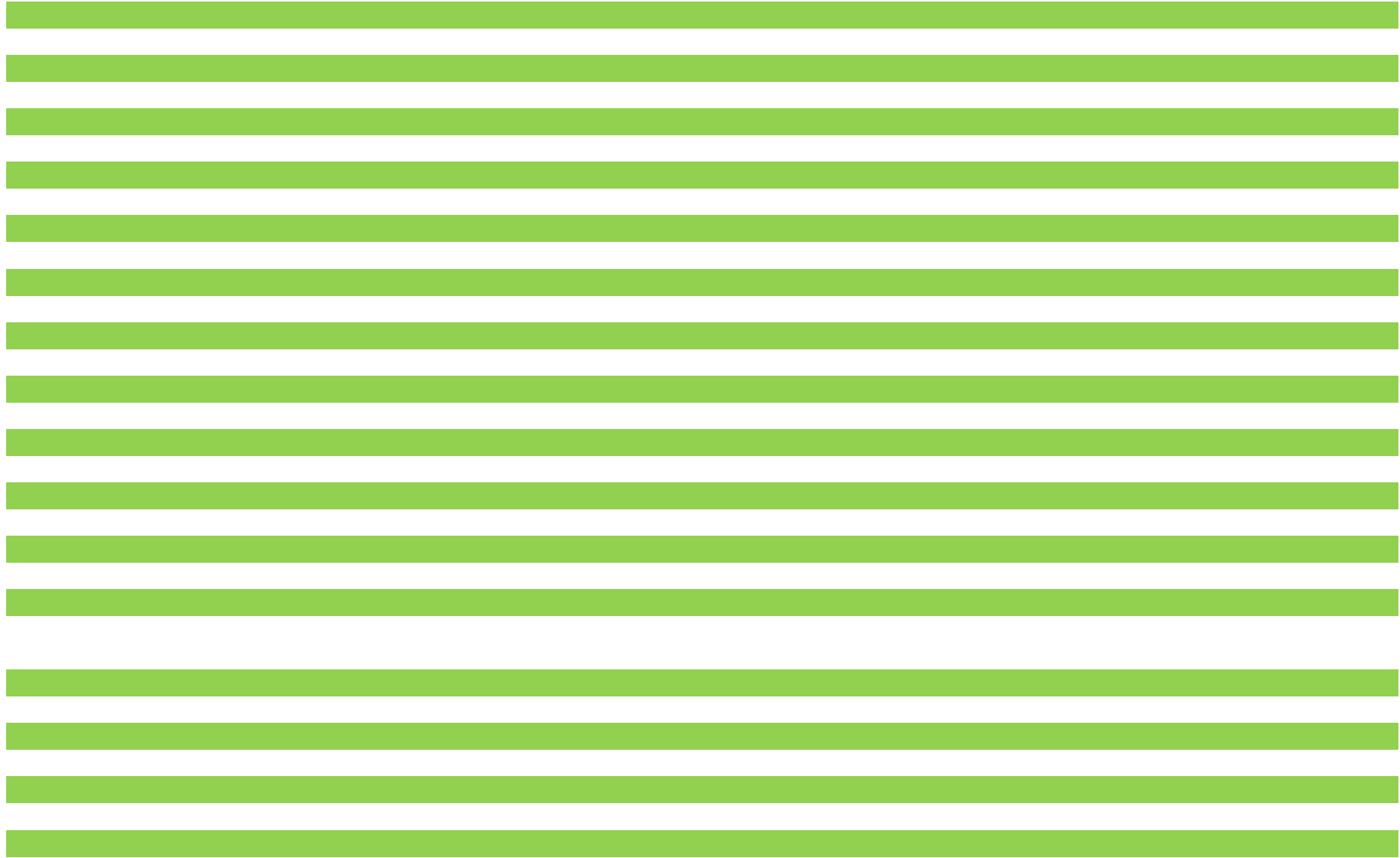
TAB]



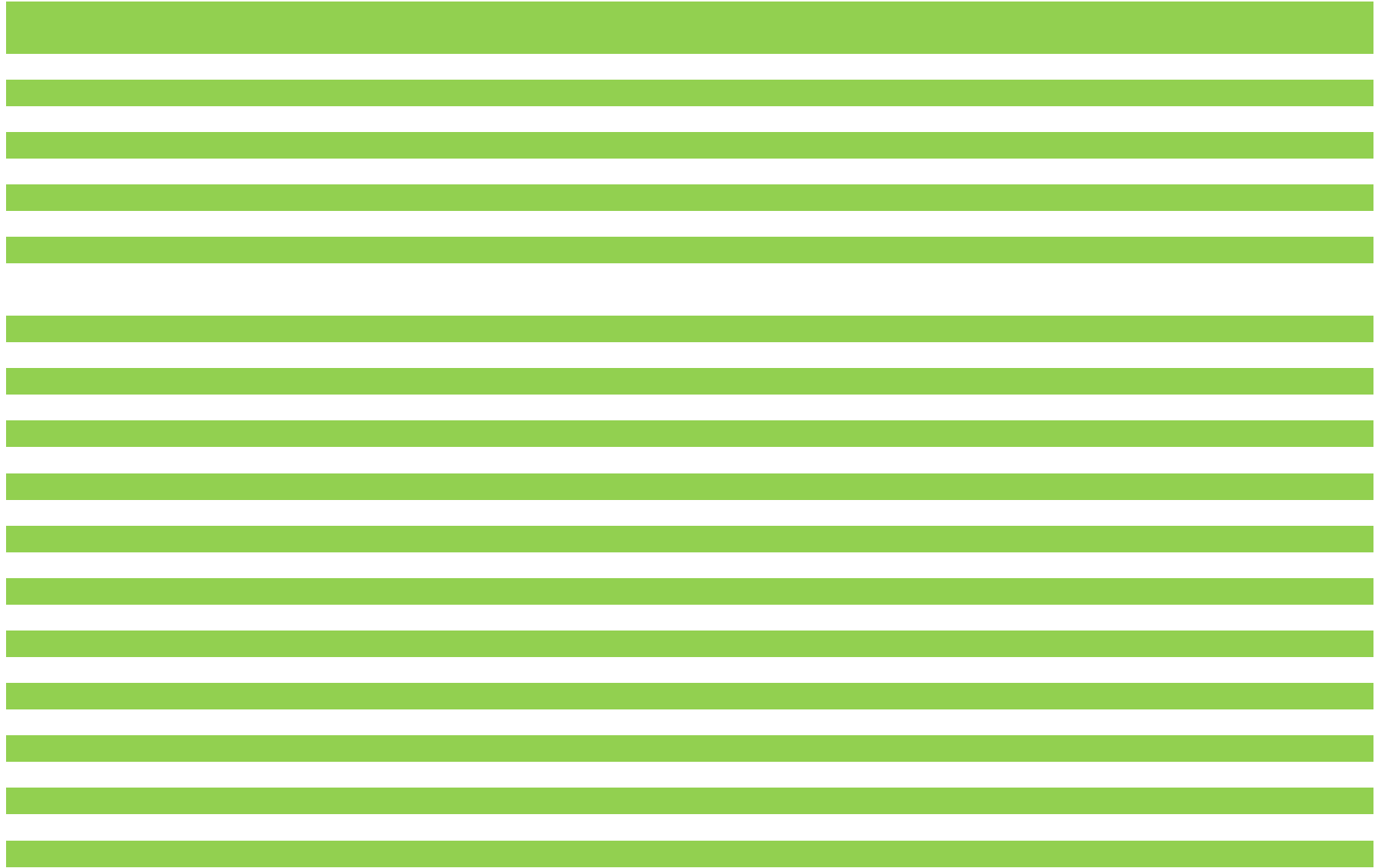
TAB]



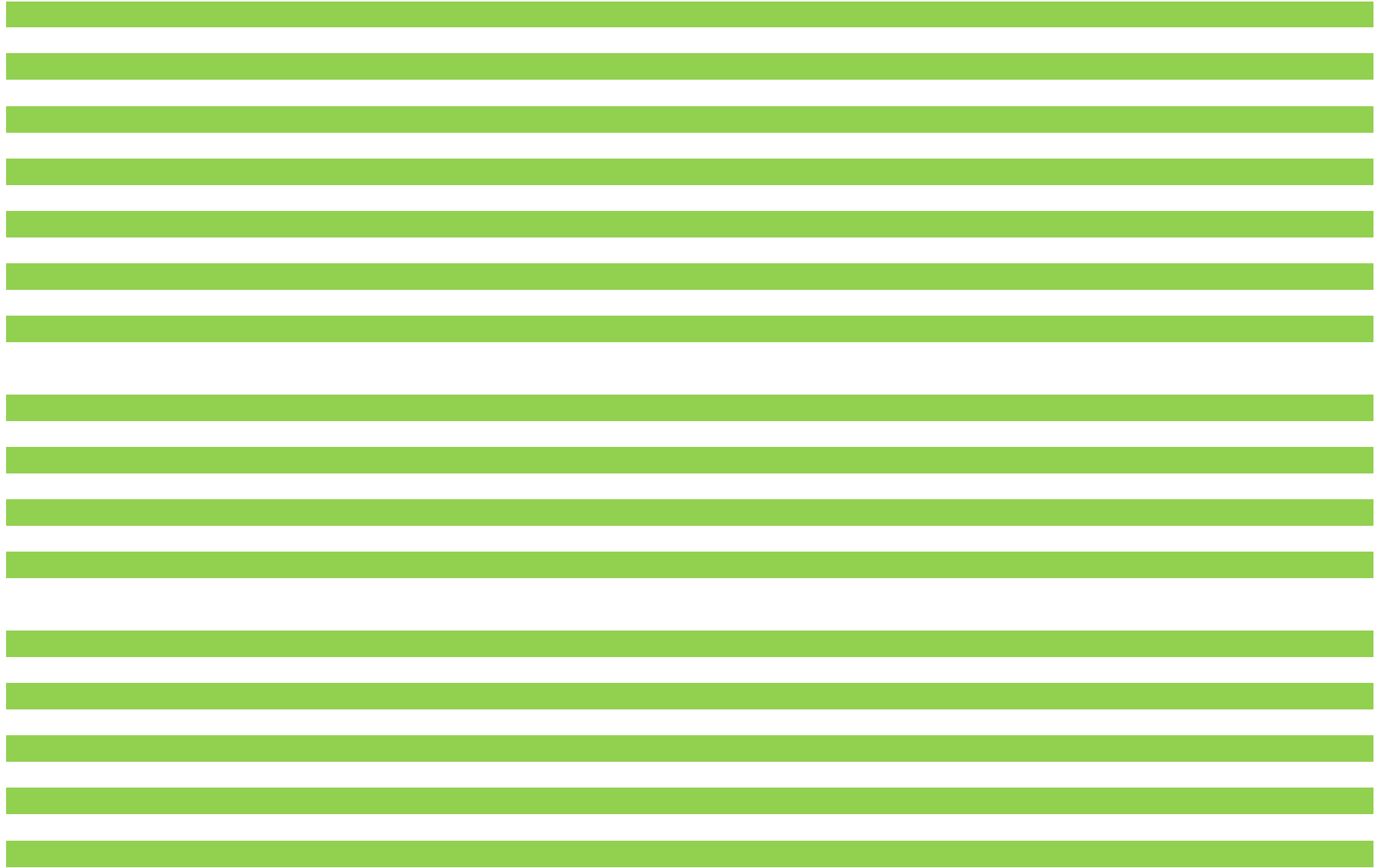
TAB]



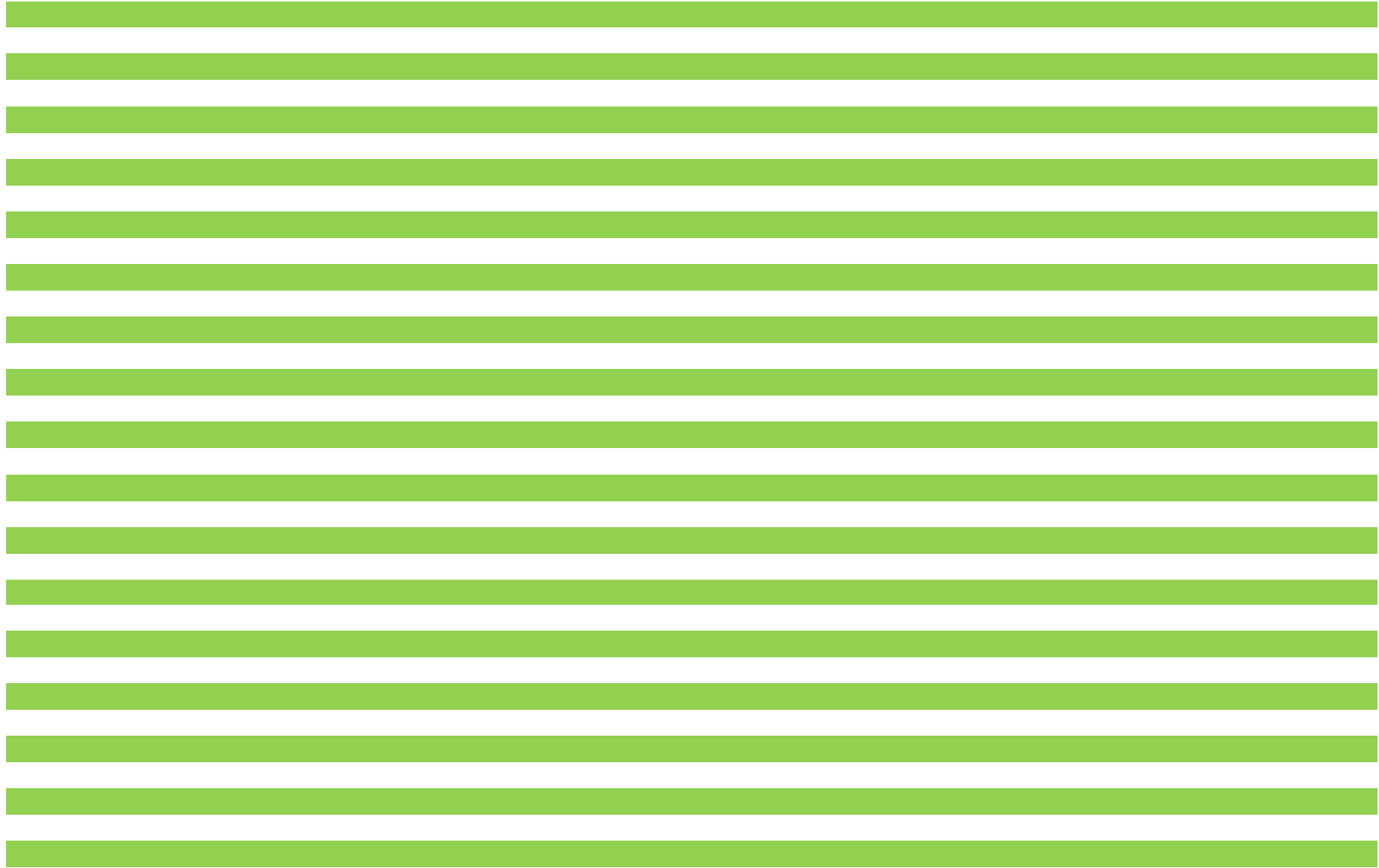
TAB]



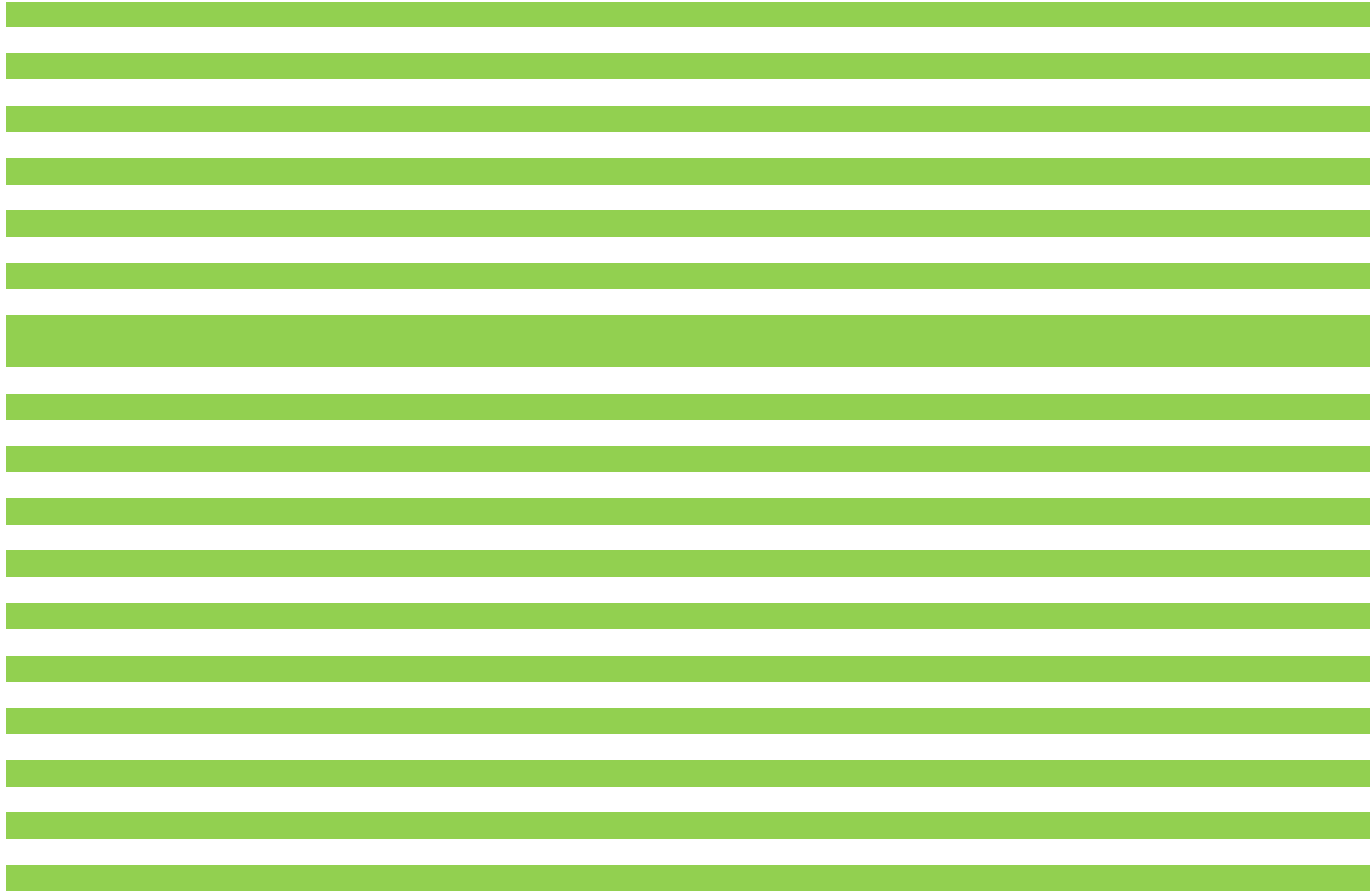
TAB]



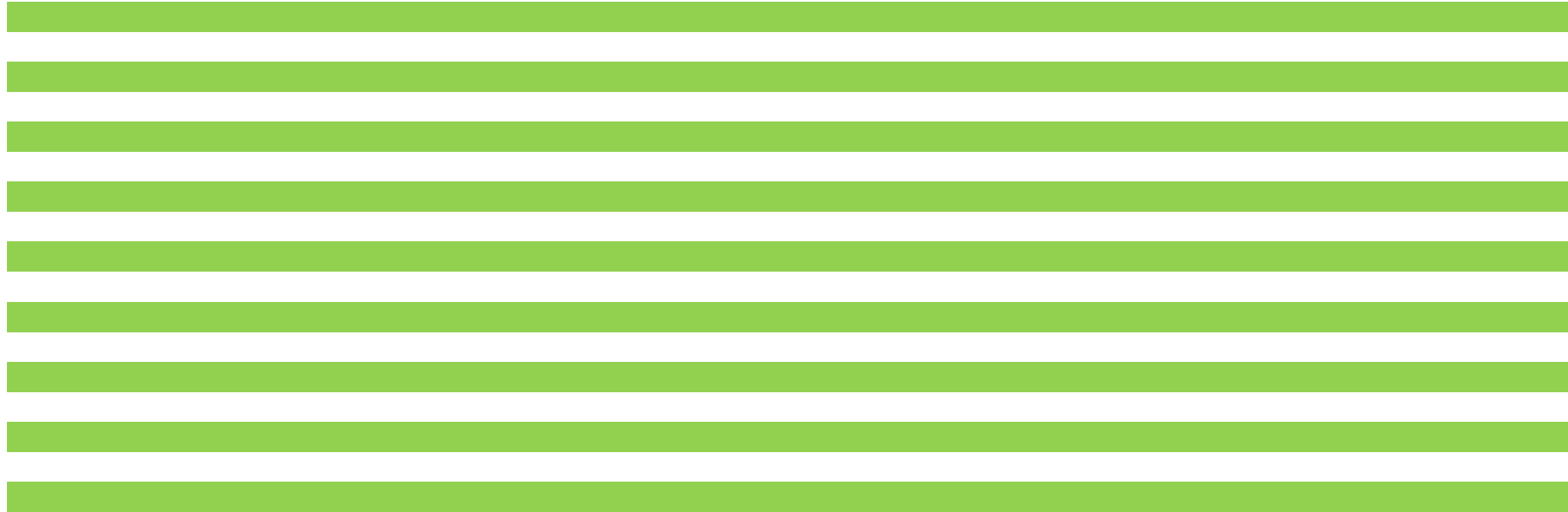
TAB]



TAB]



TAB]



TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

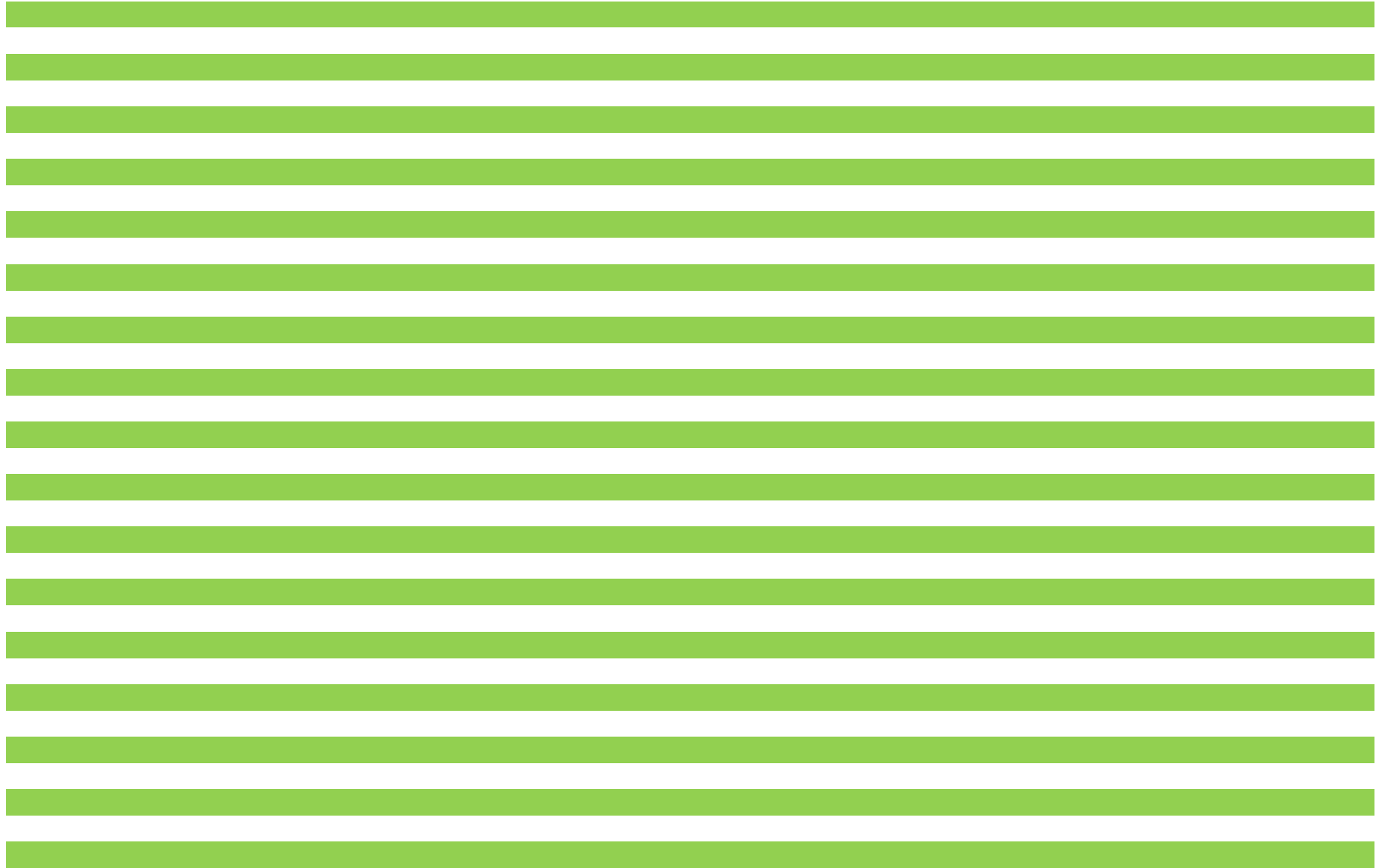
TAB]

TAB]

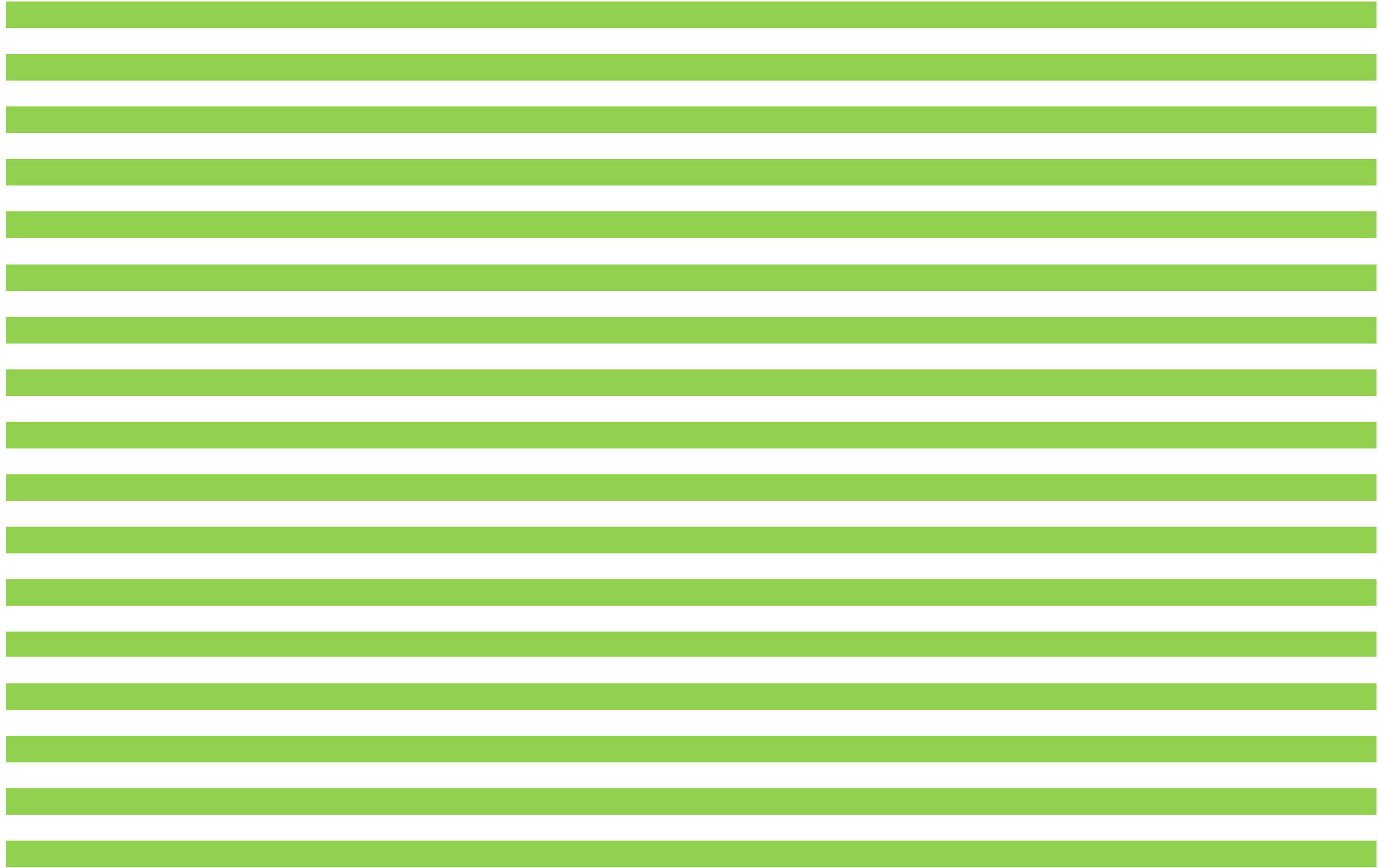
TAB]

TAB]

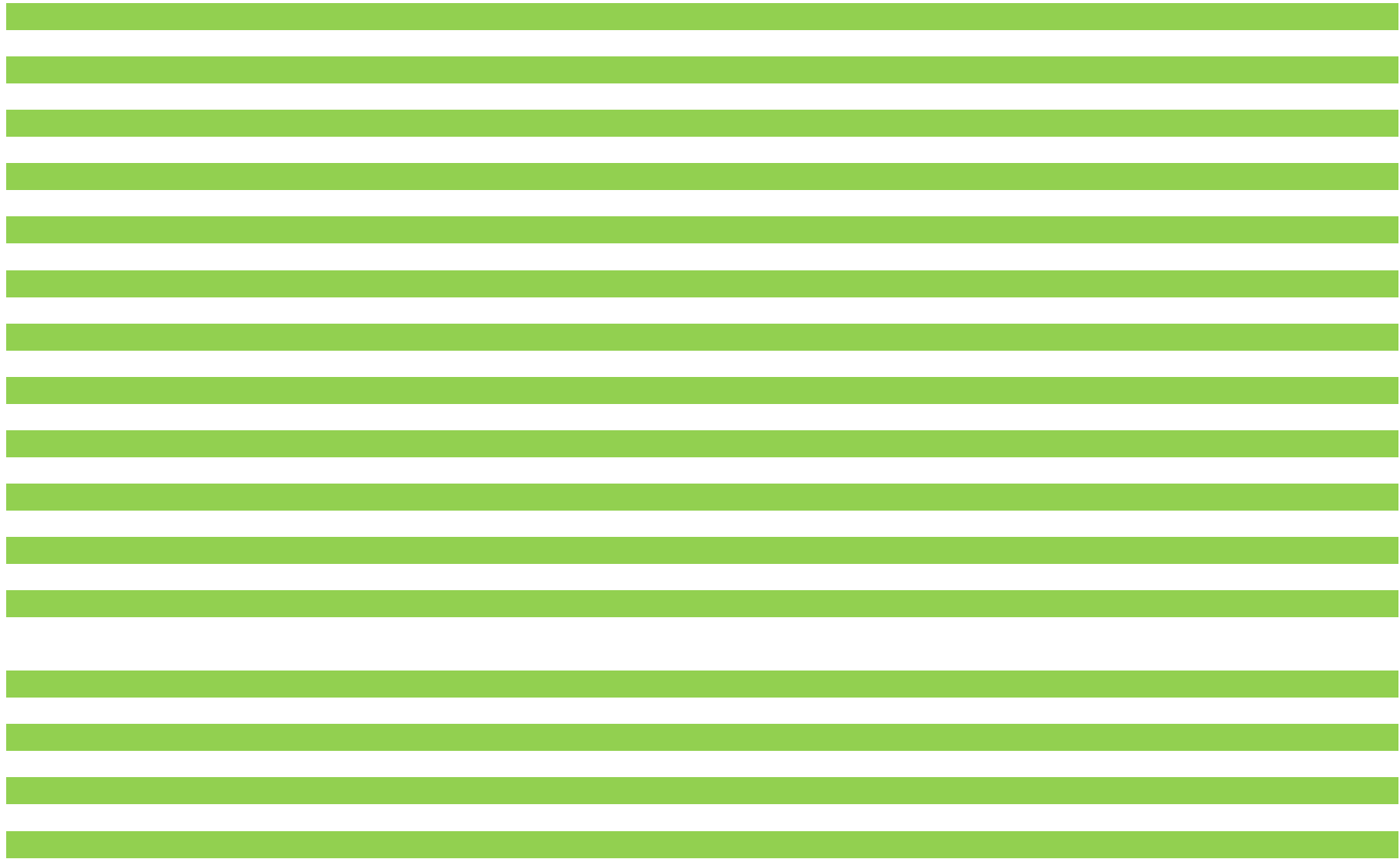
TAB]



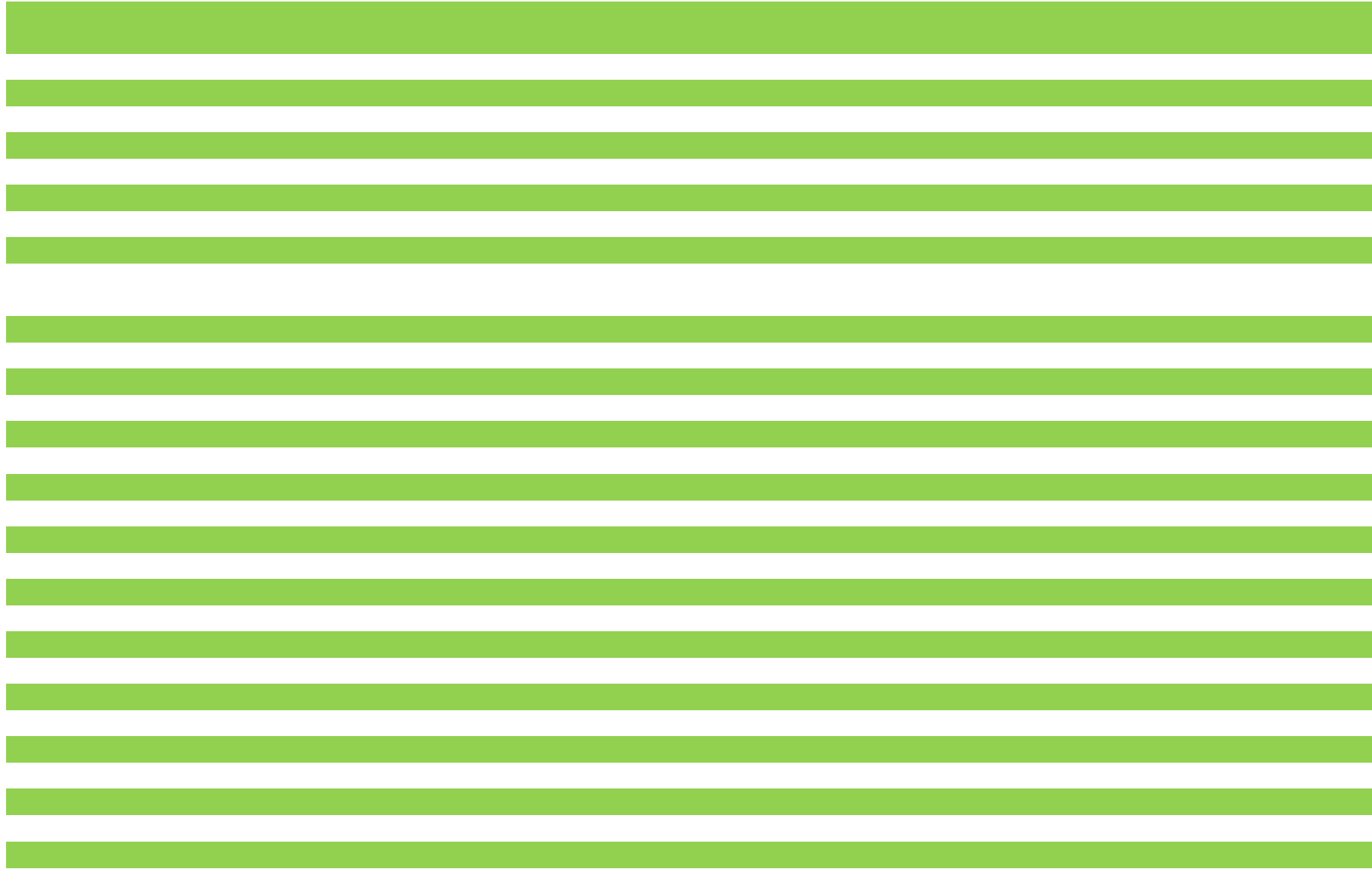
TAB]



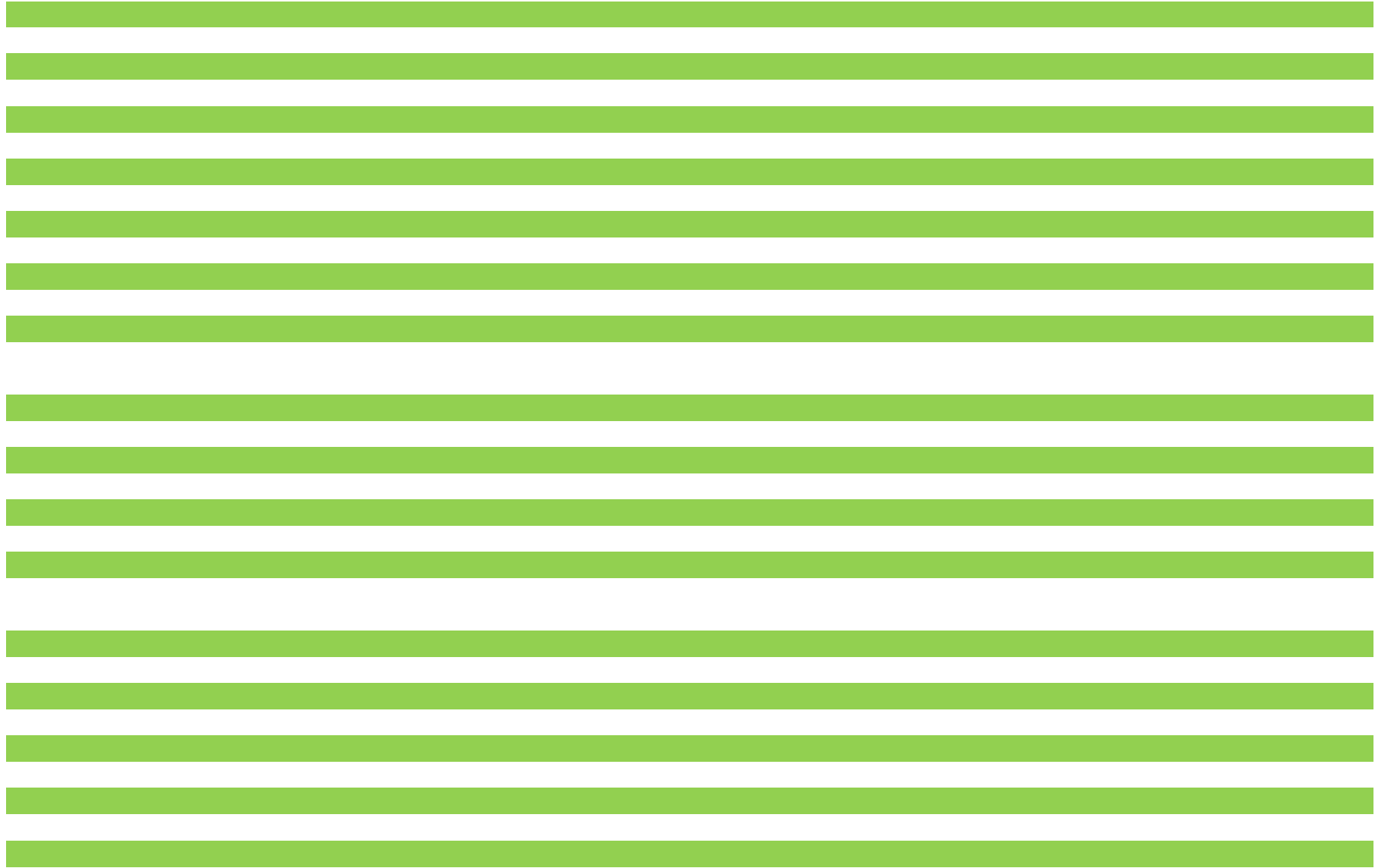
TAB]



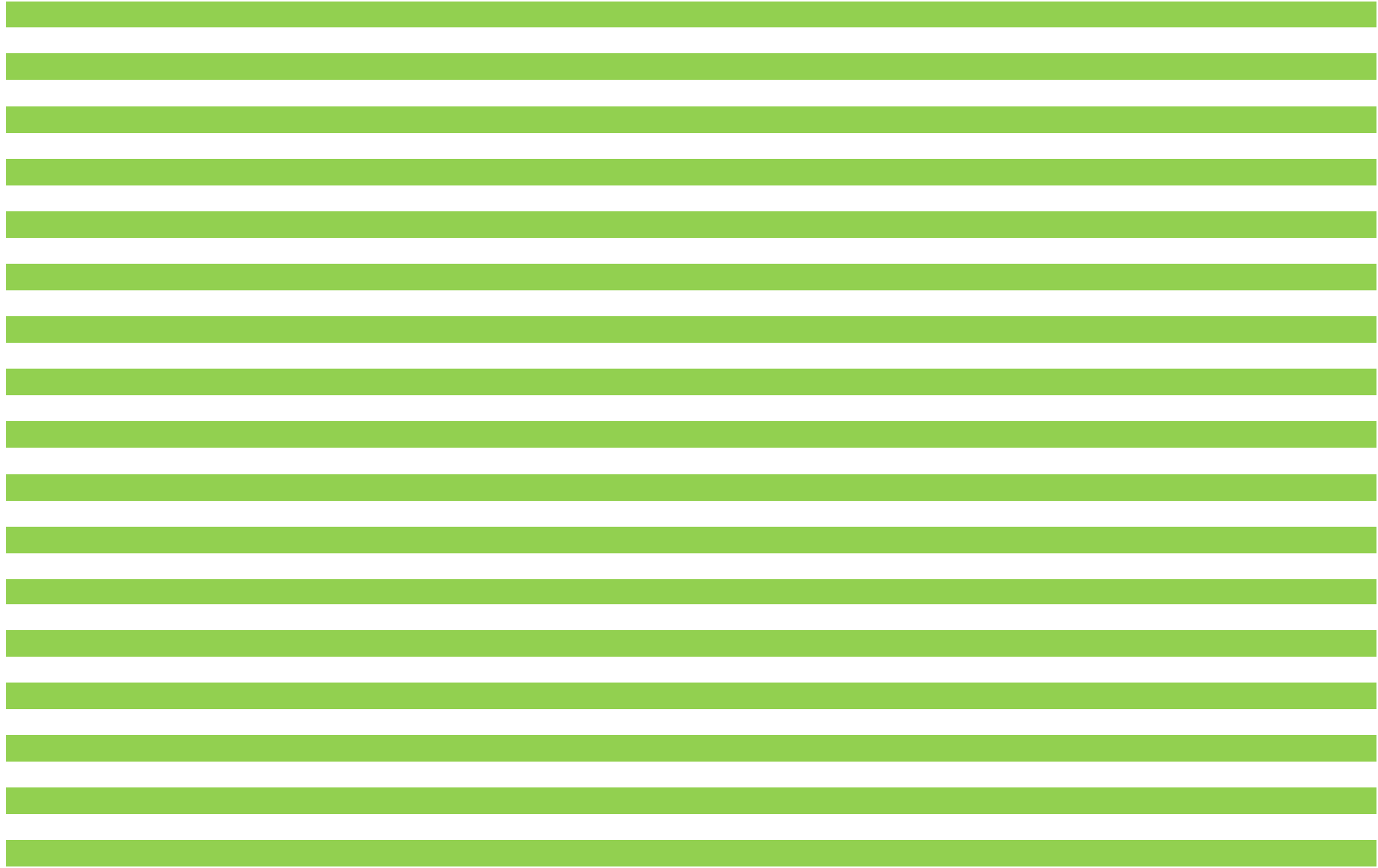
TAB]



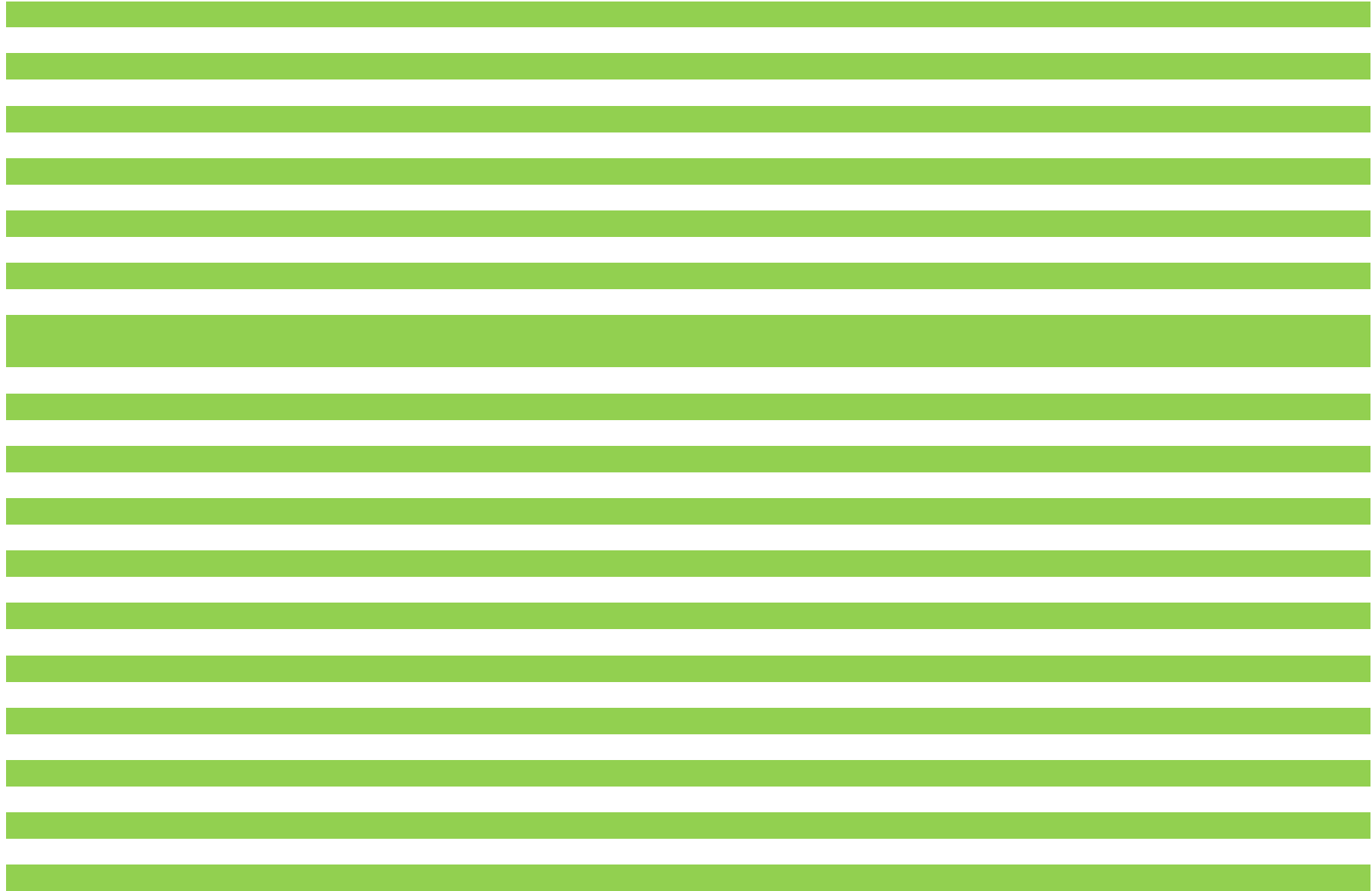
TAB]



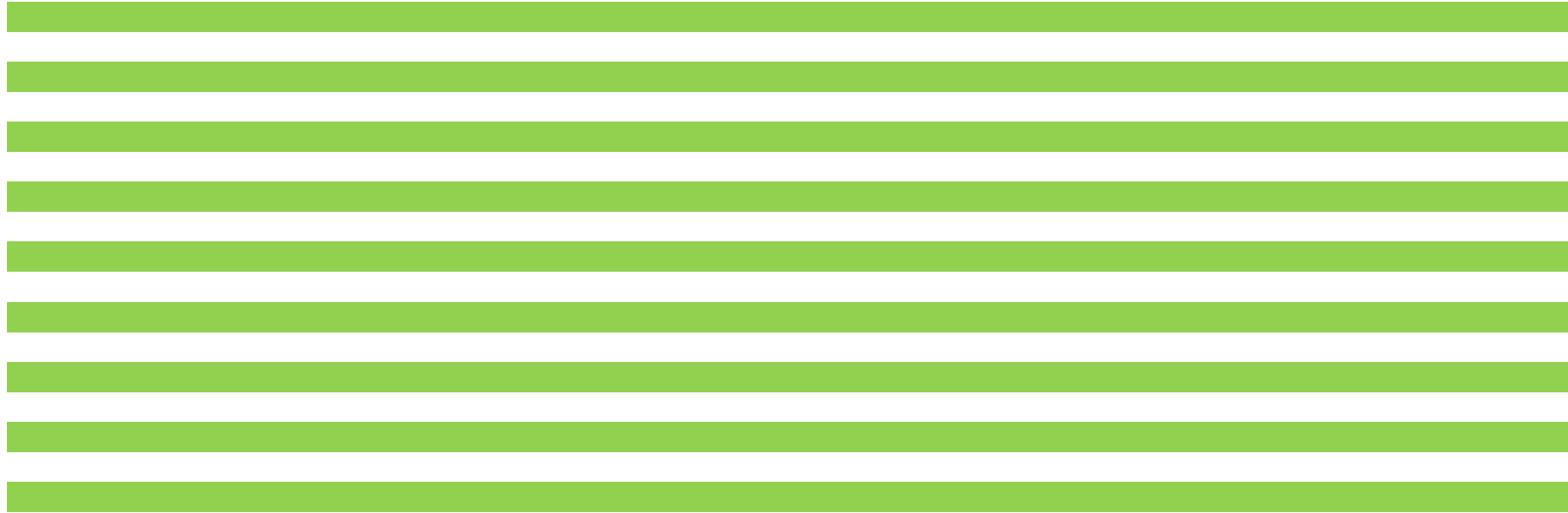
TAB]



TAB]



TAB]



TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

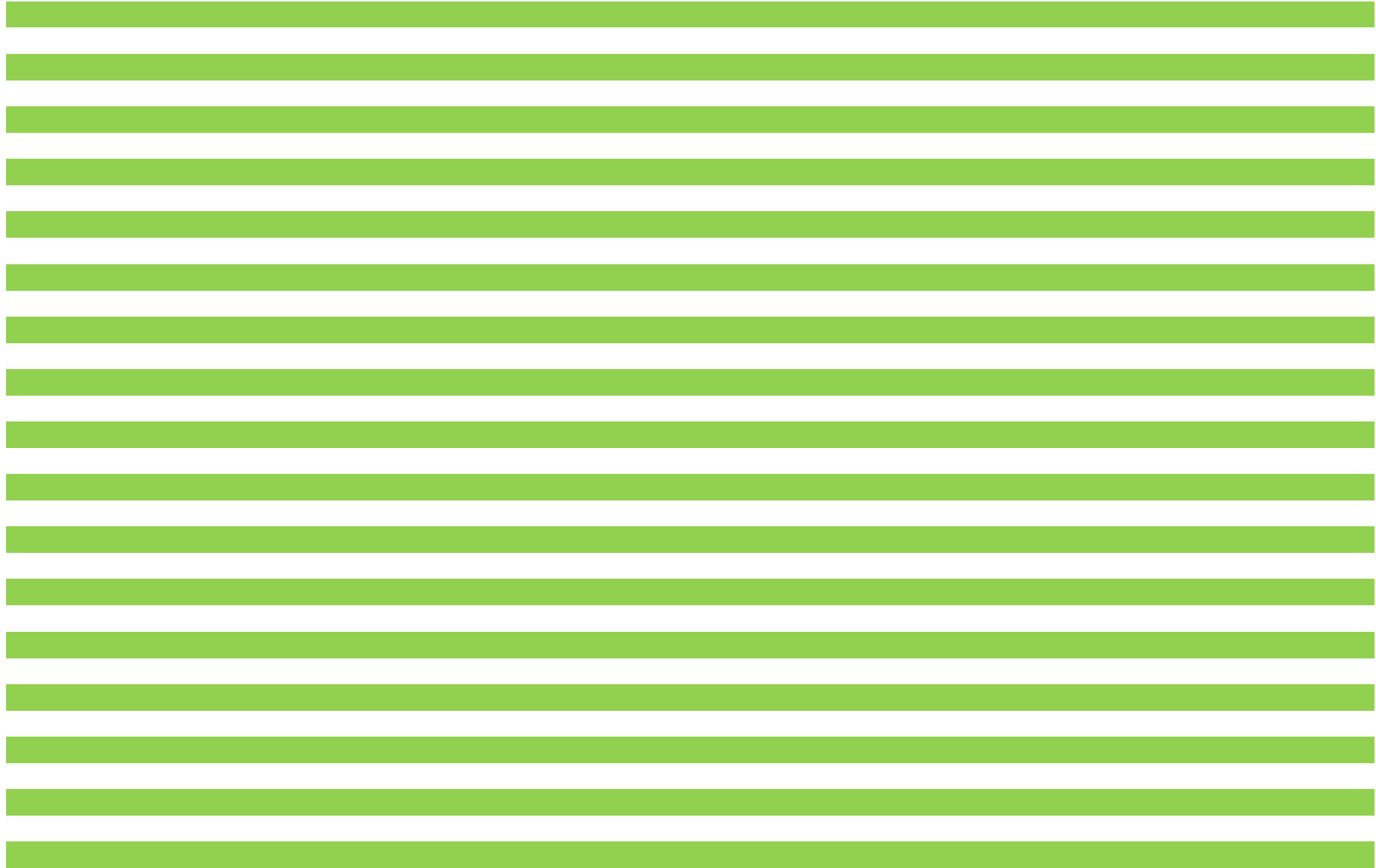
TAB]

TAB]

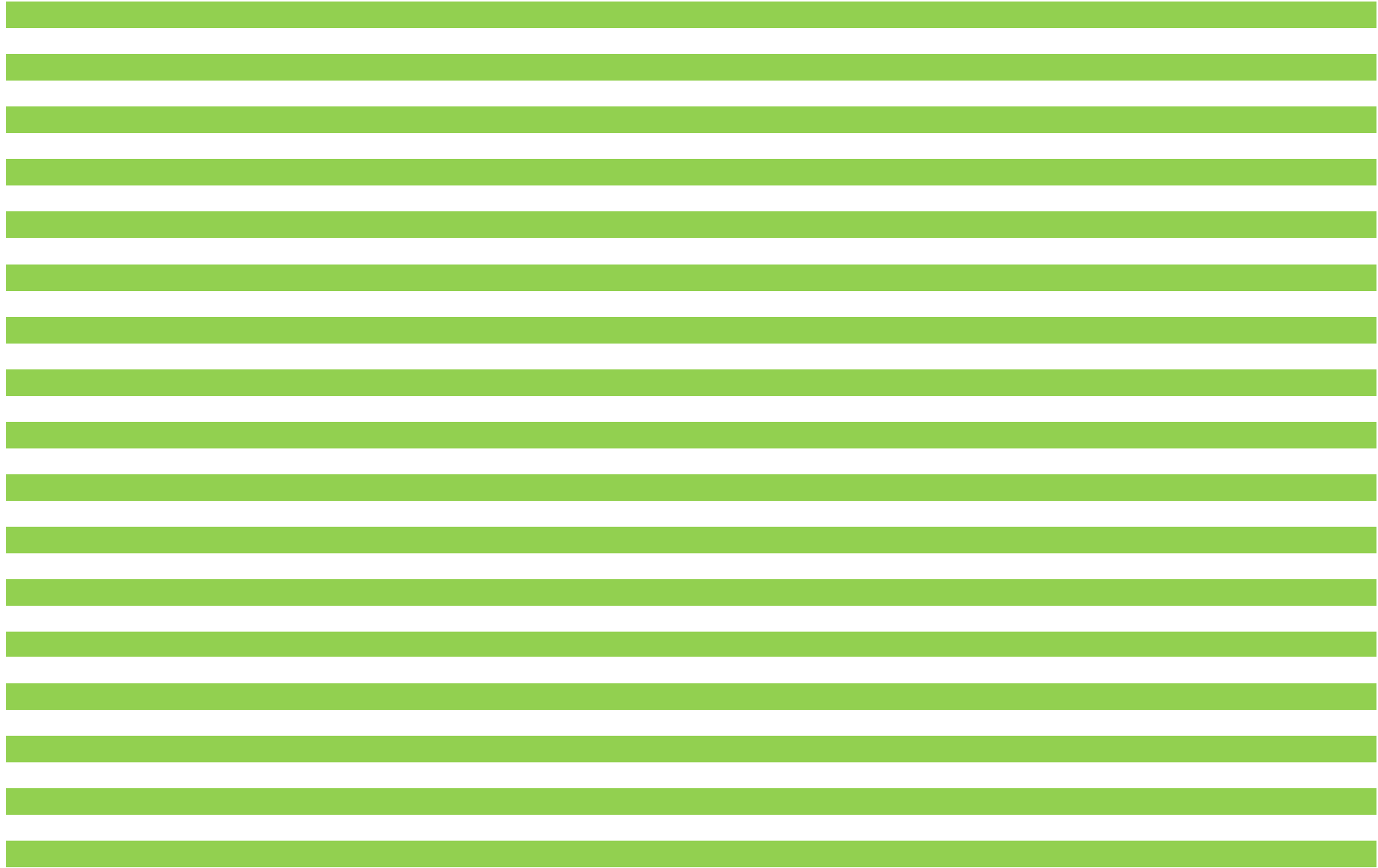
TAB]

TAB]

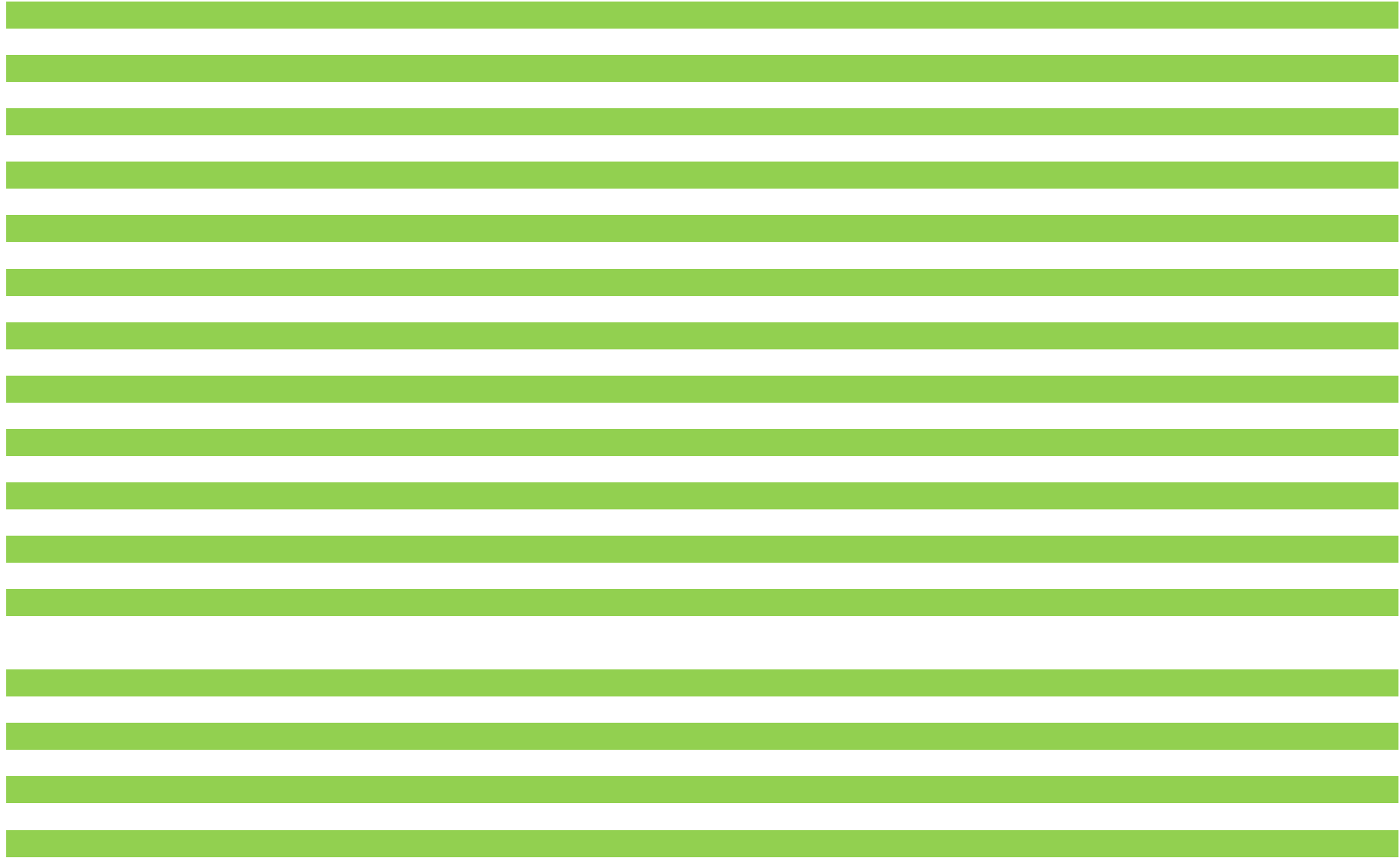
TAB]



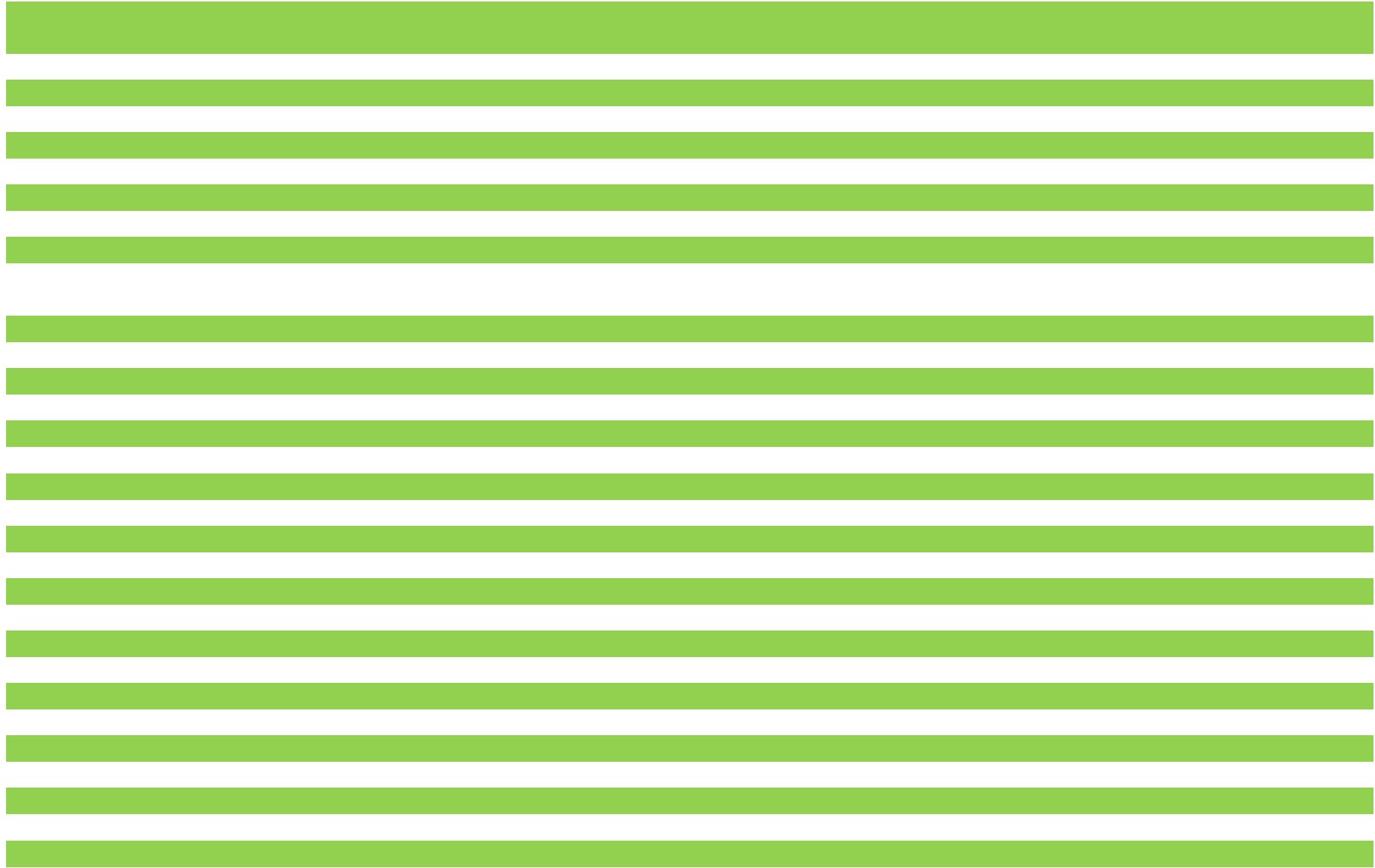
TAB]



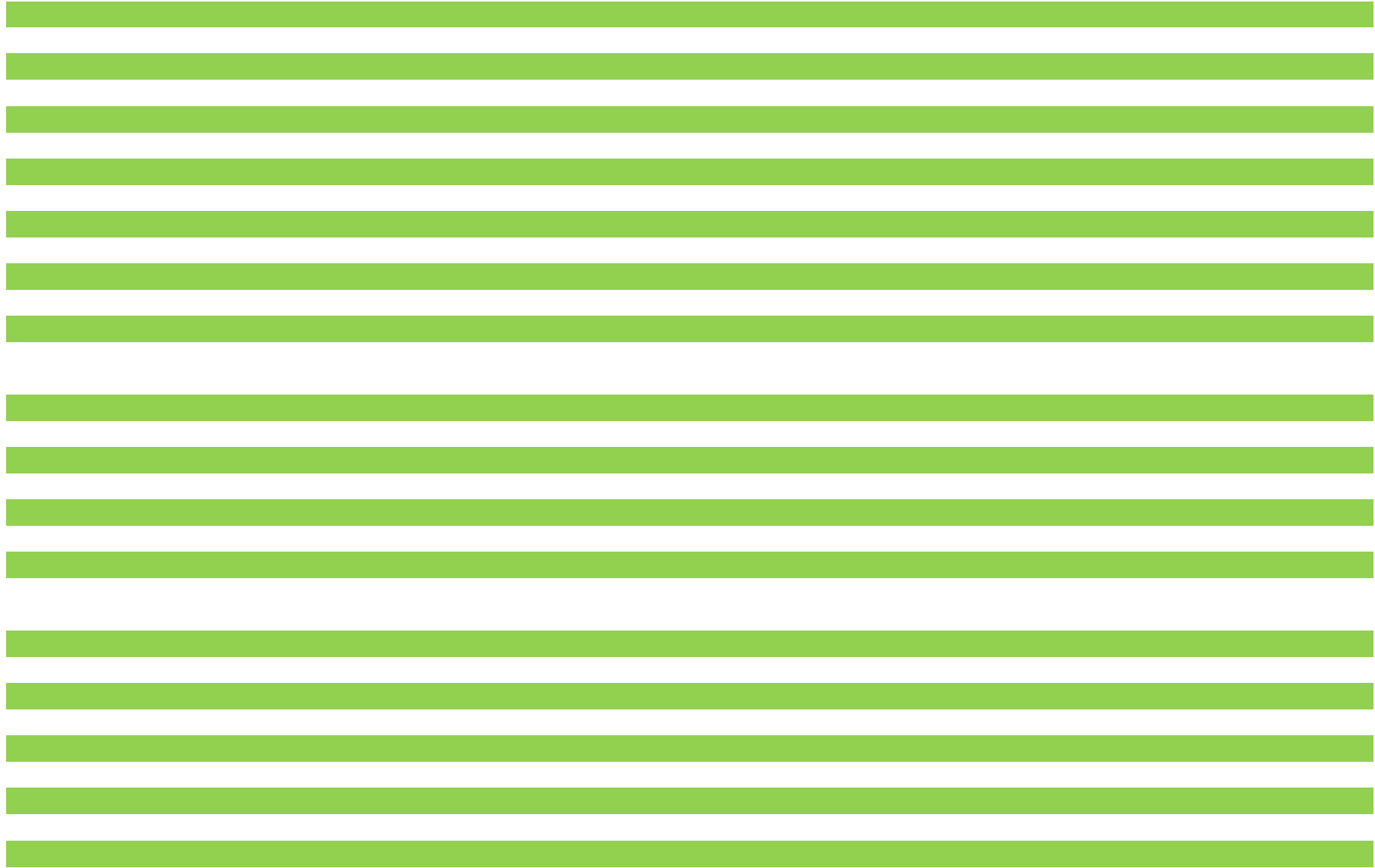
TAB]



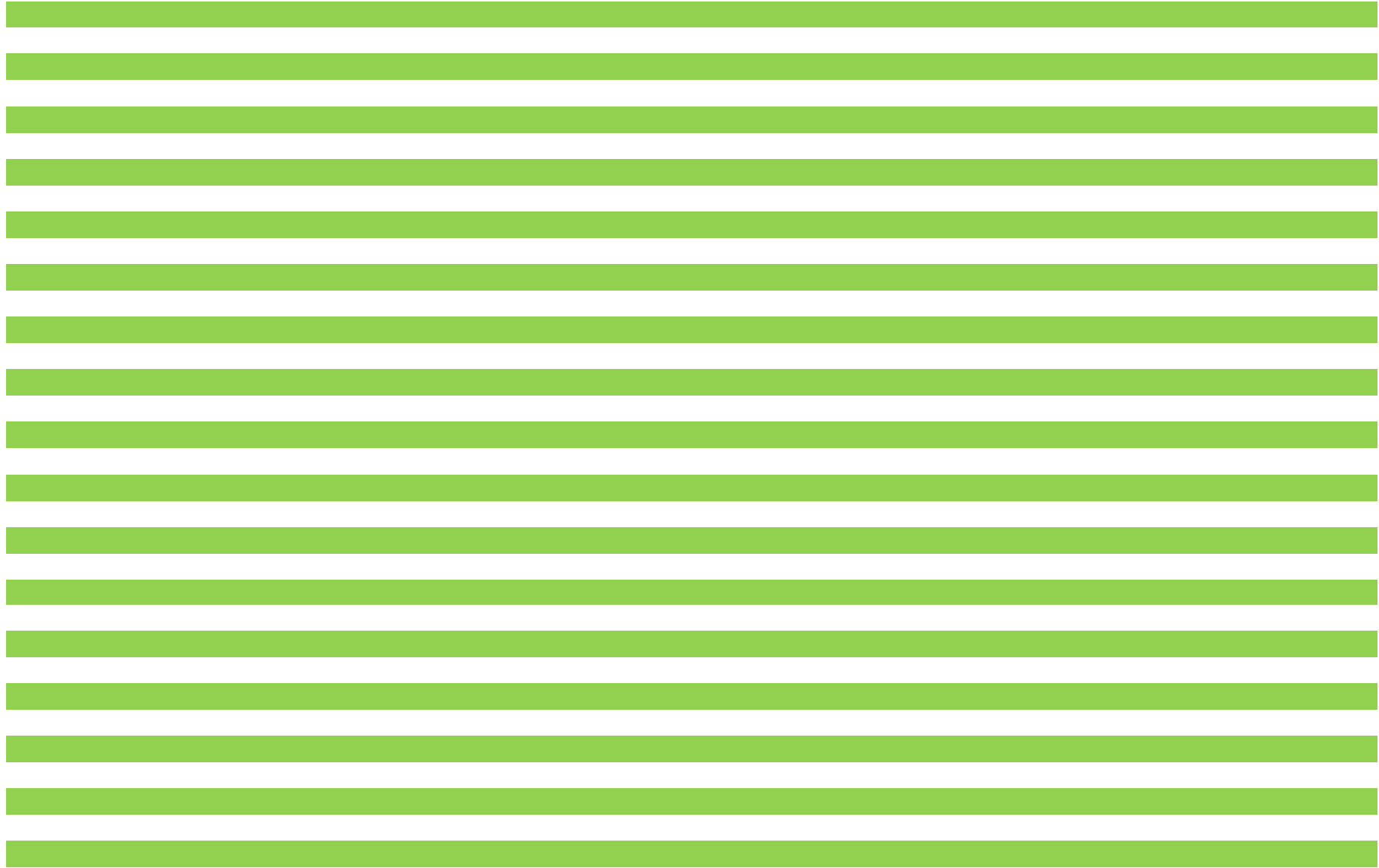
TAB]



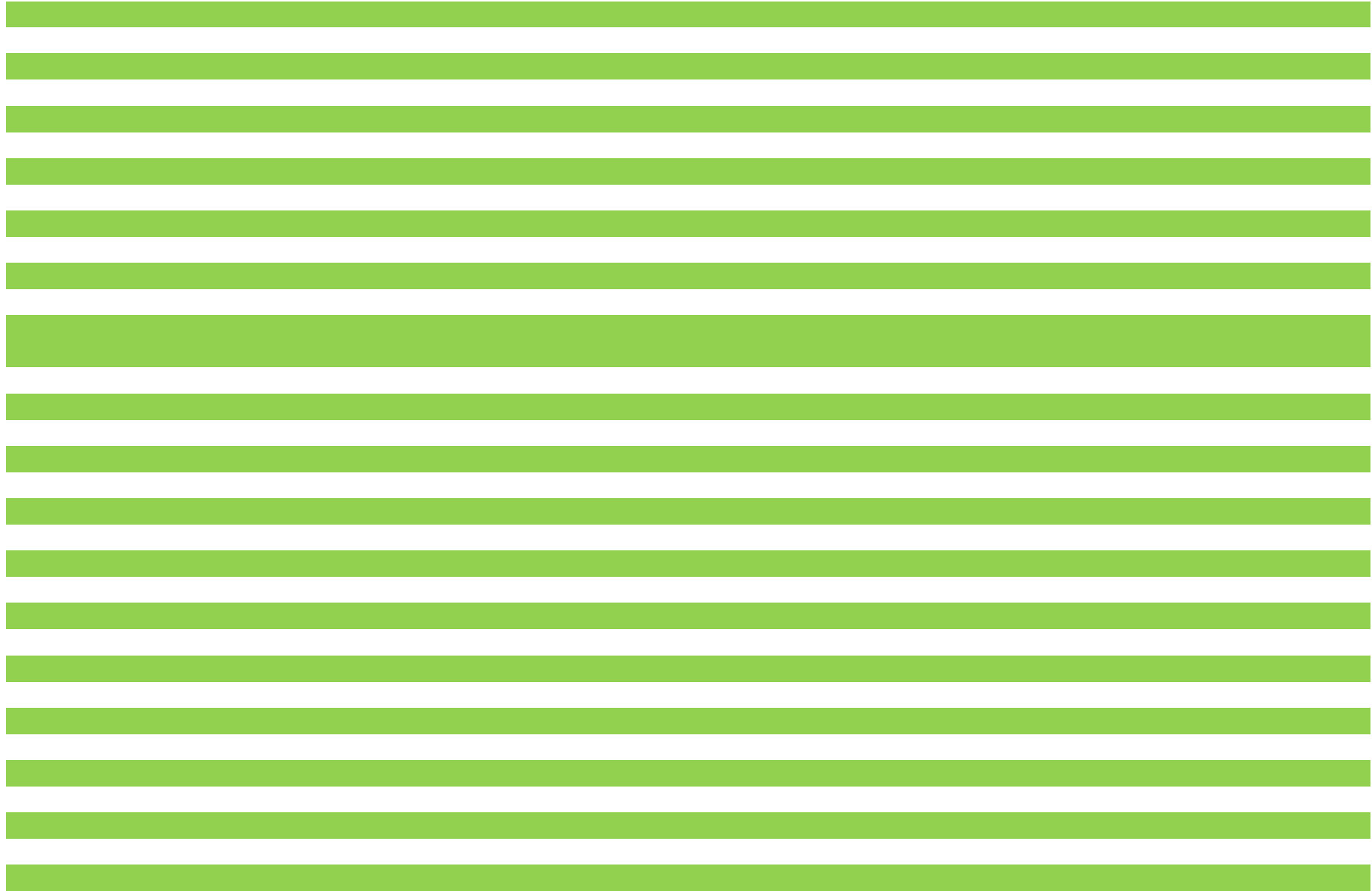
TAB]



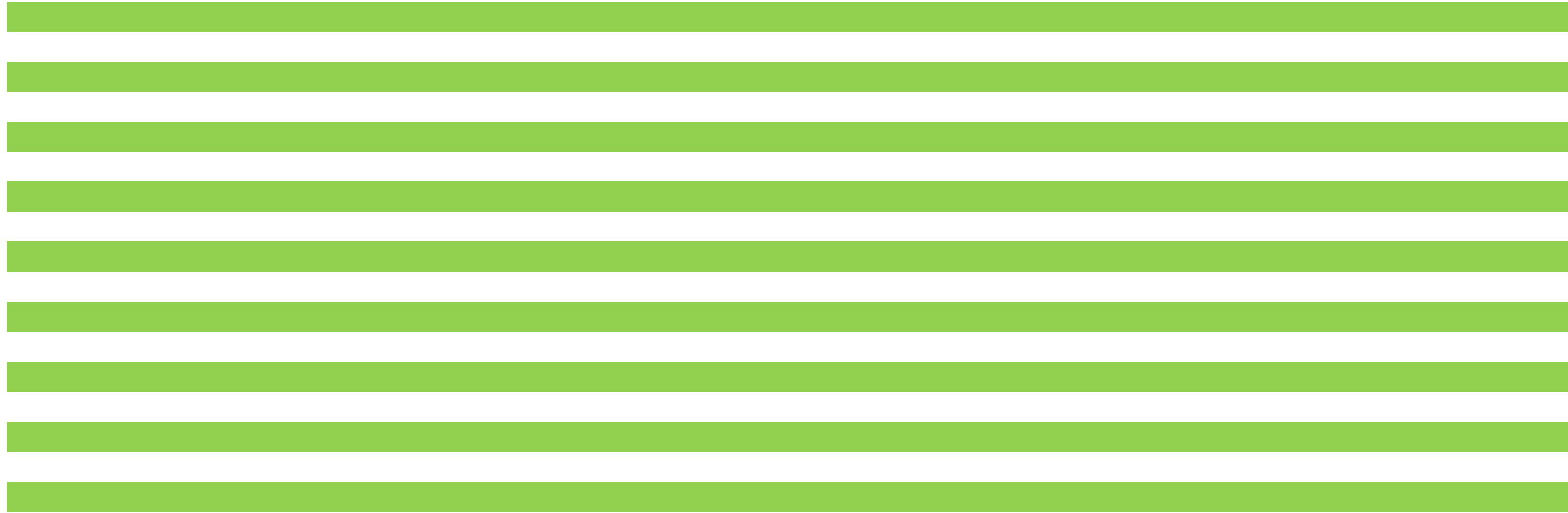
TAB]



TAB]



TAB]



TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

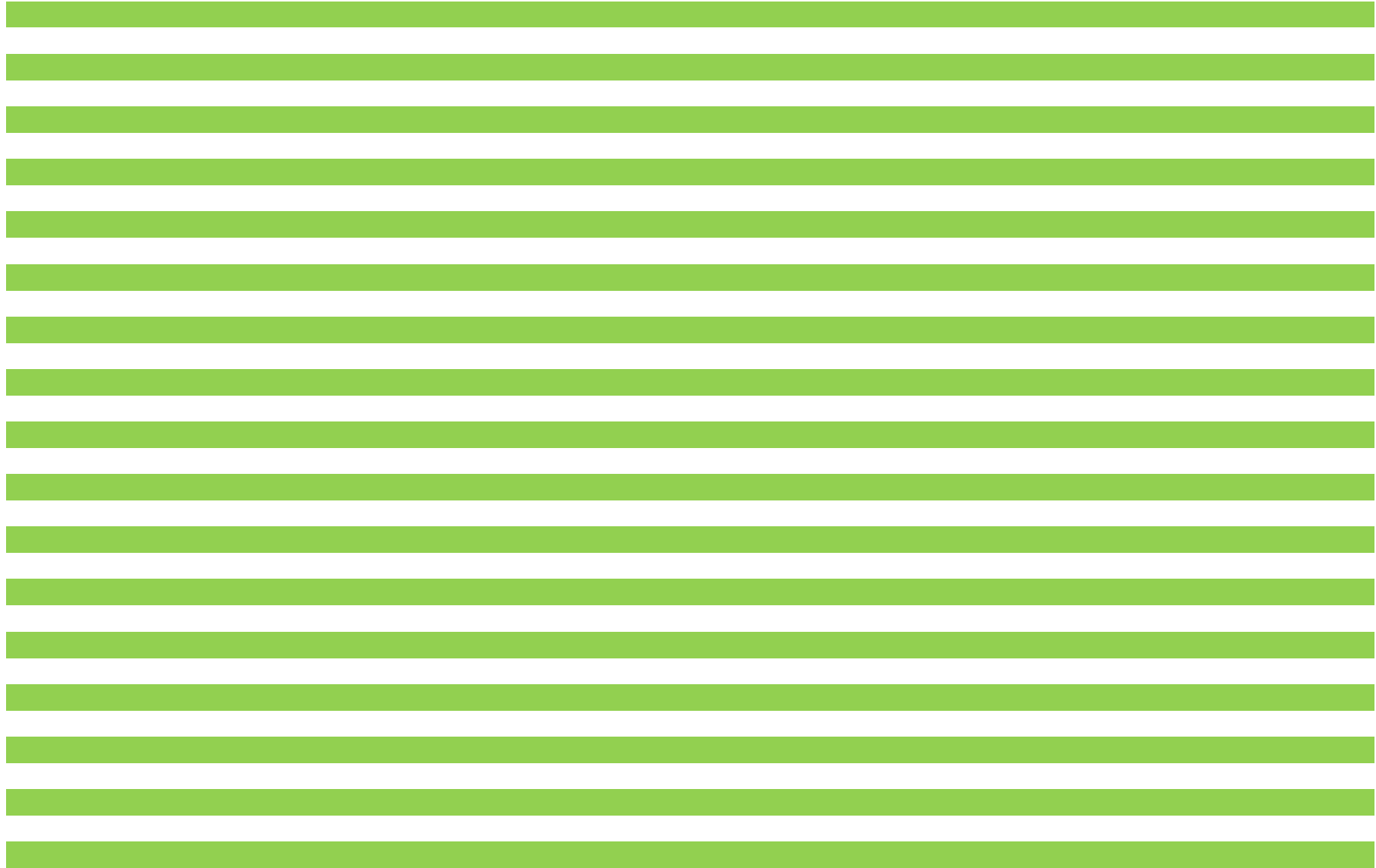
TAB]

TAB]

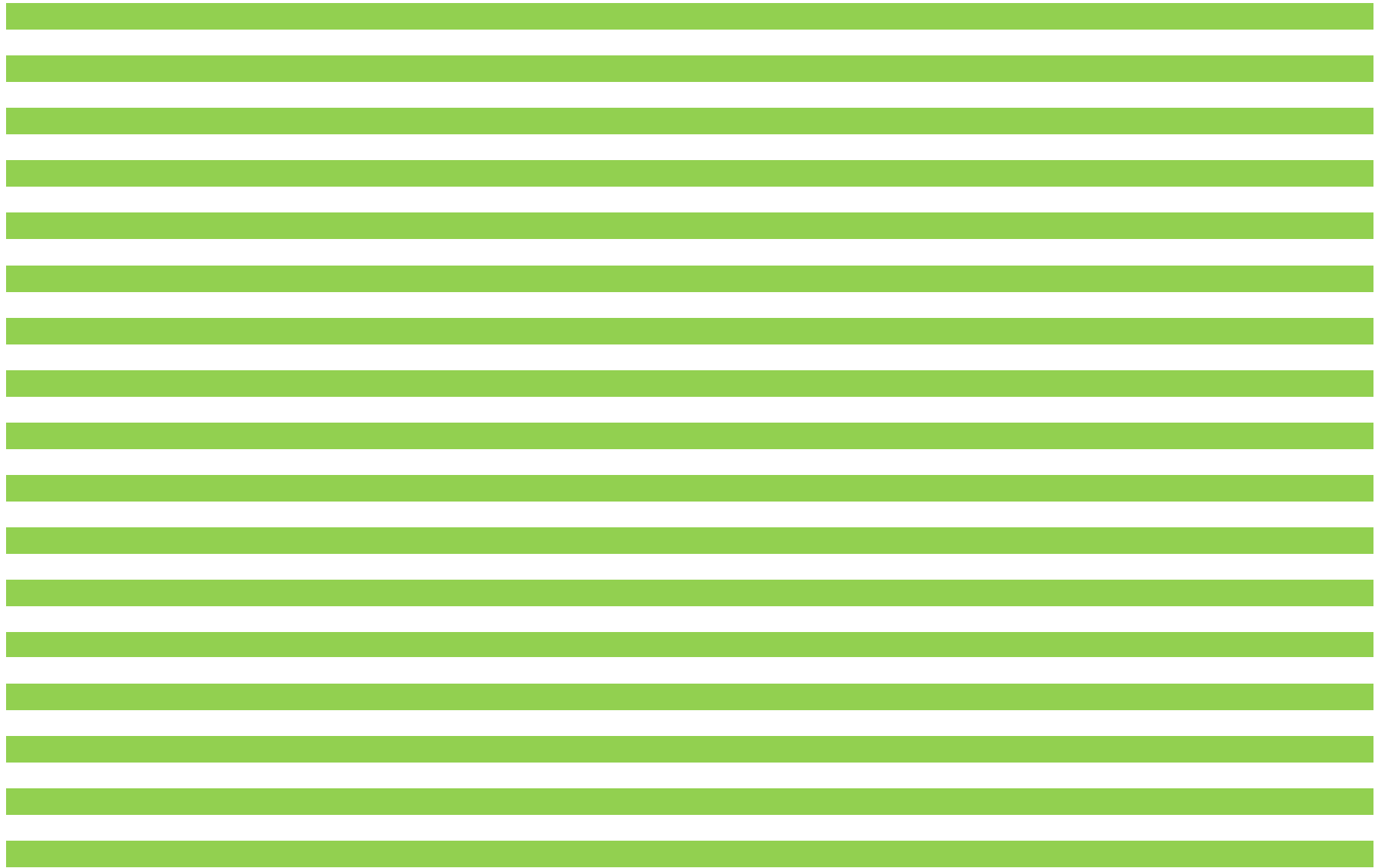
TAB]

TAB]

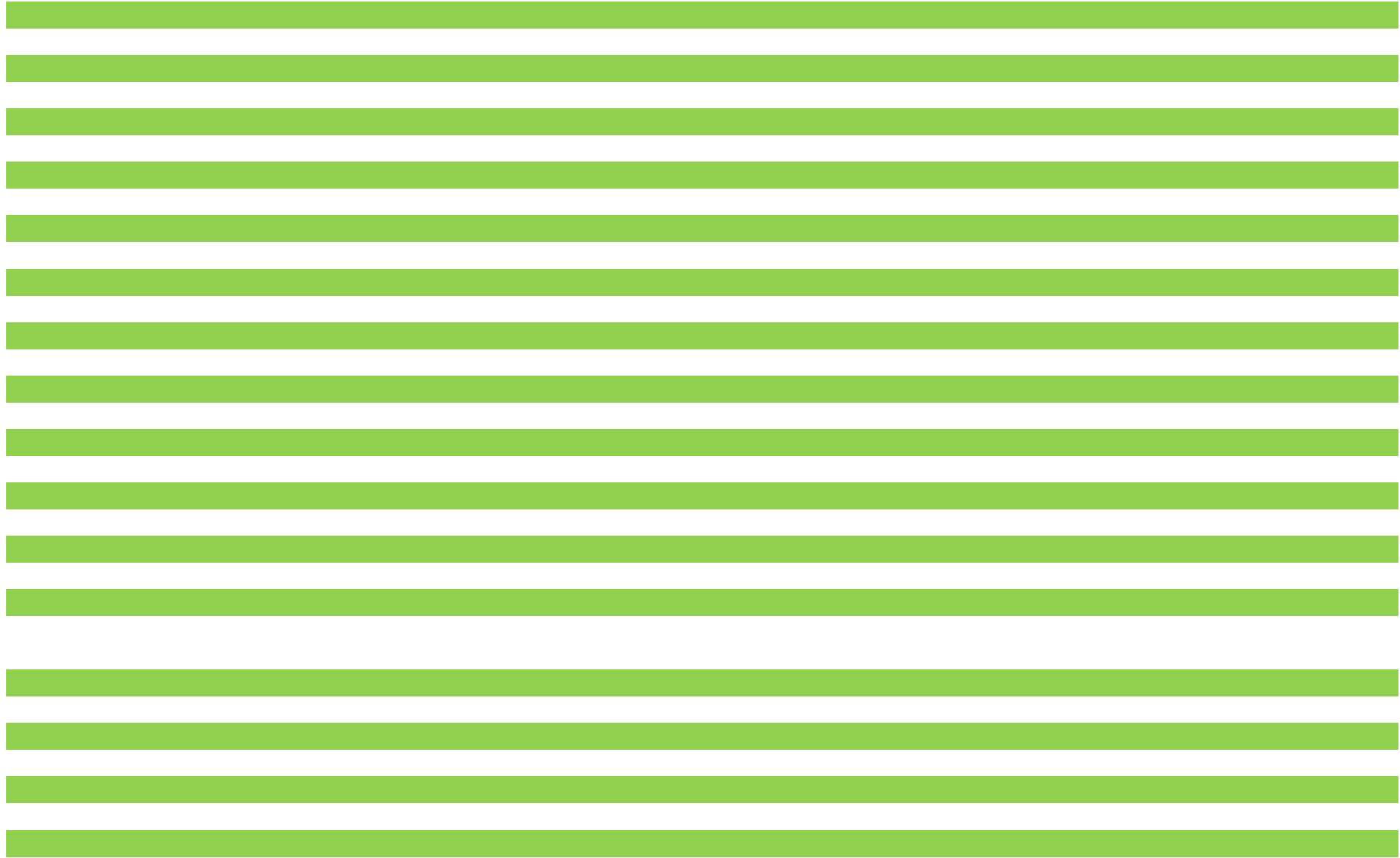
TAB]



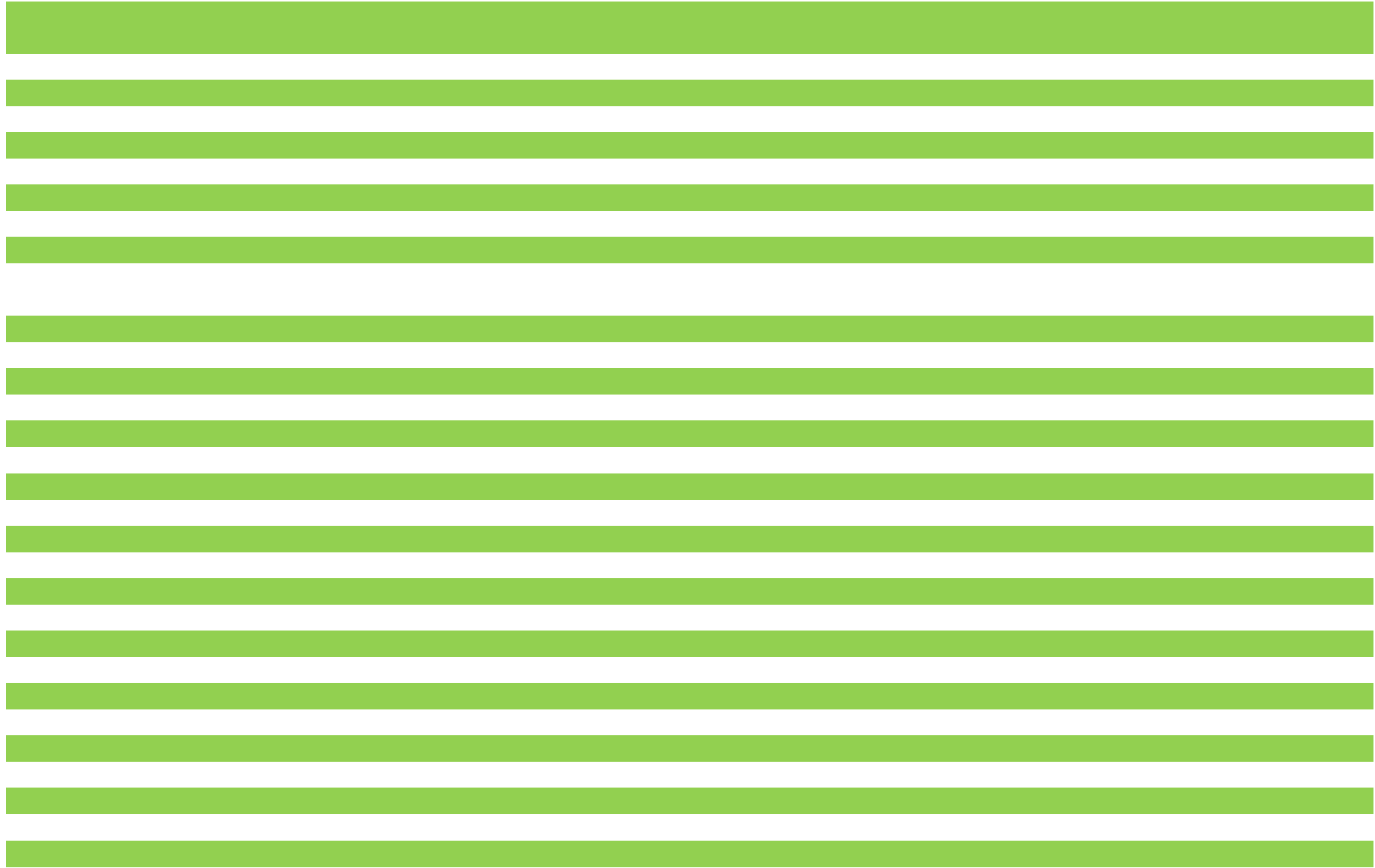
TAB]



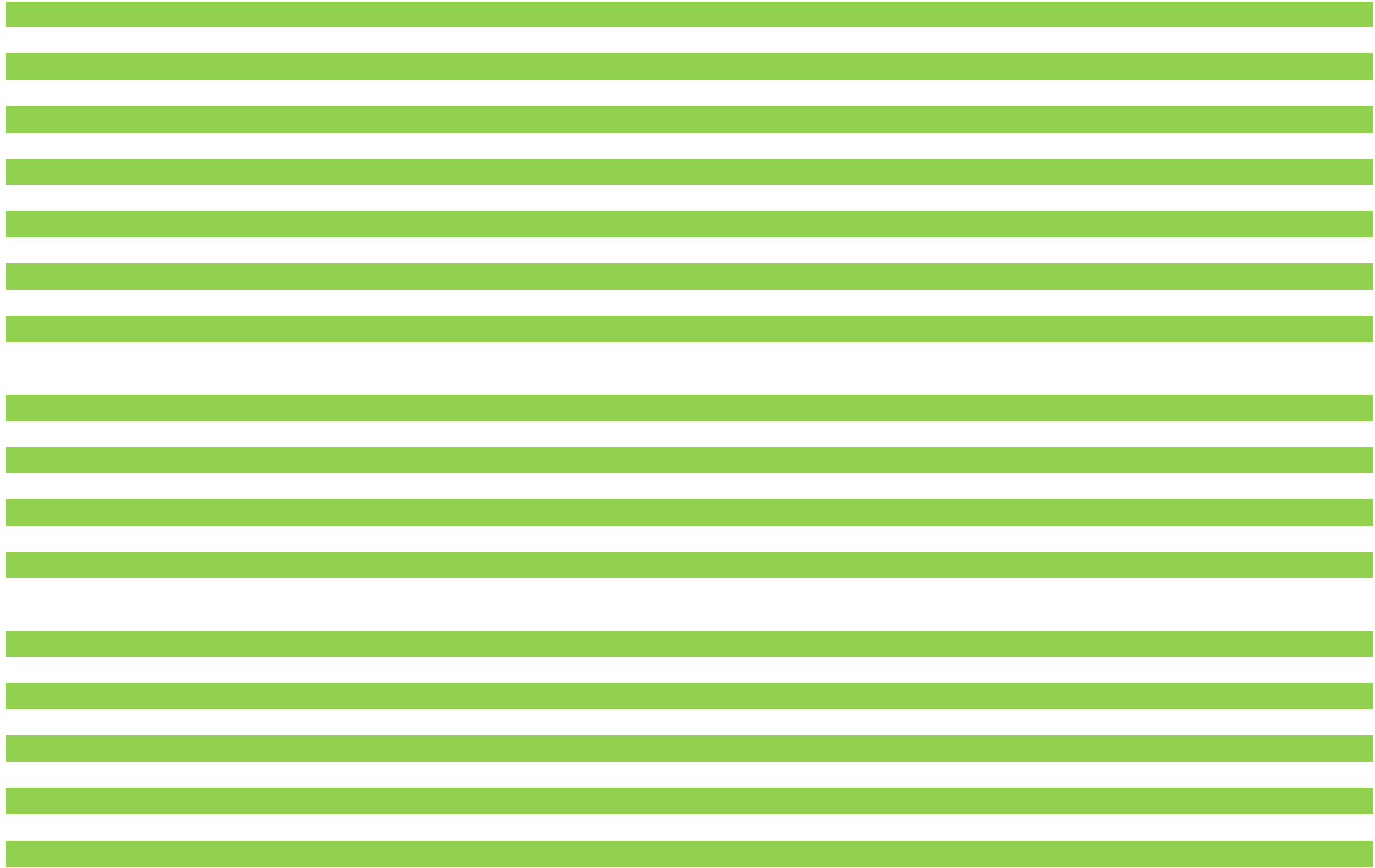
TAB]



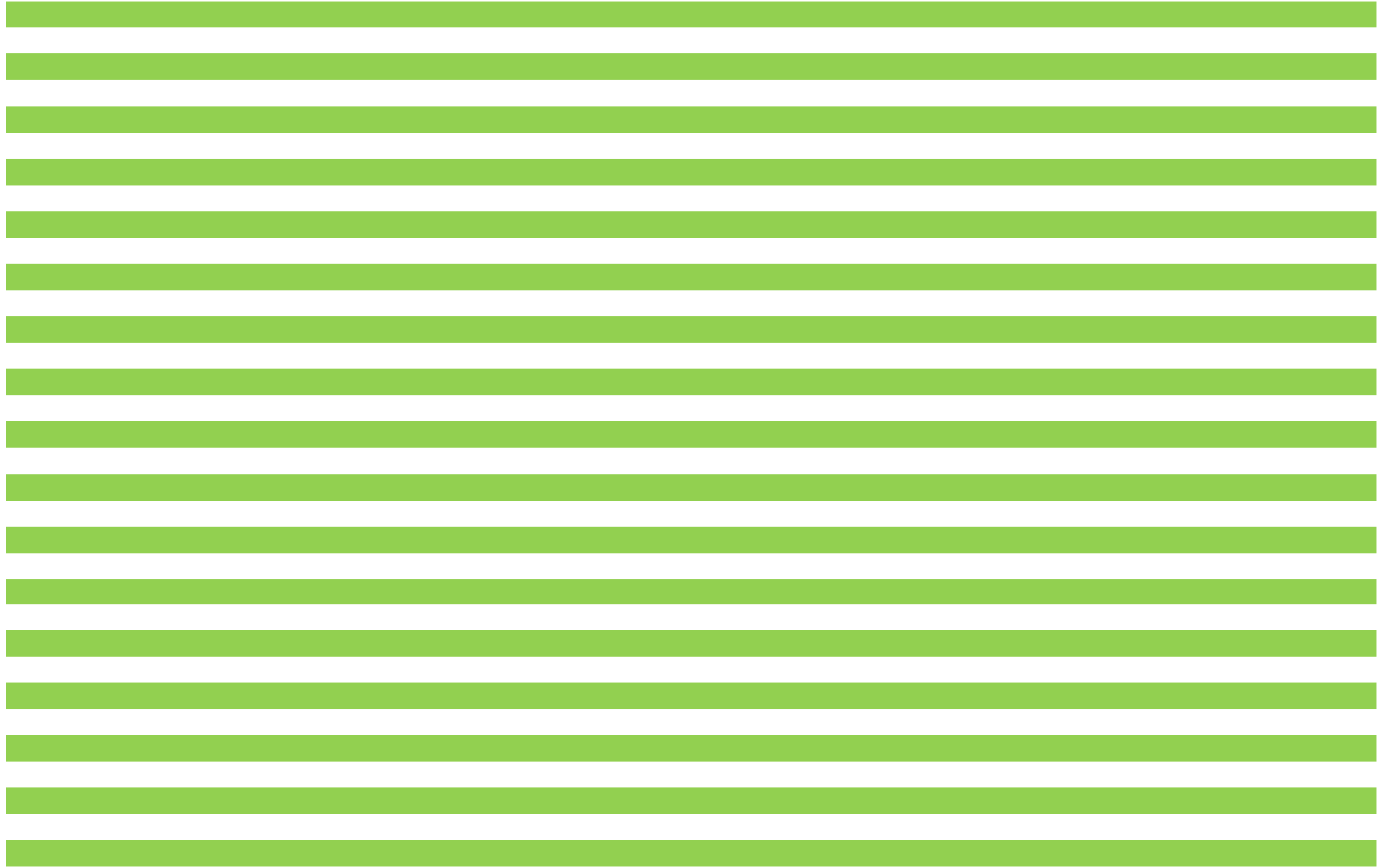
TAB]



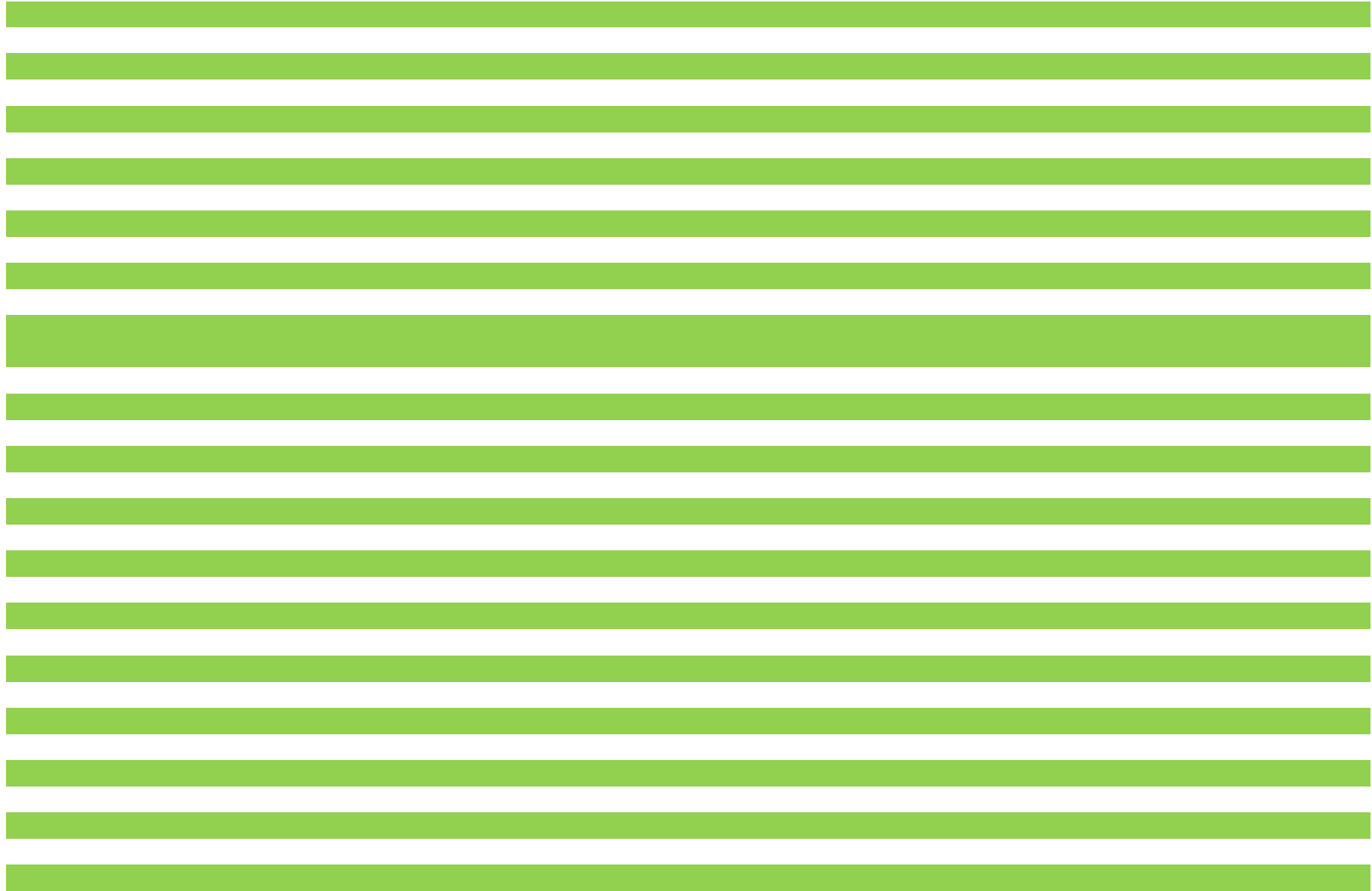
TAB]



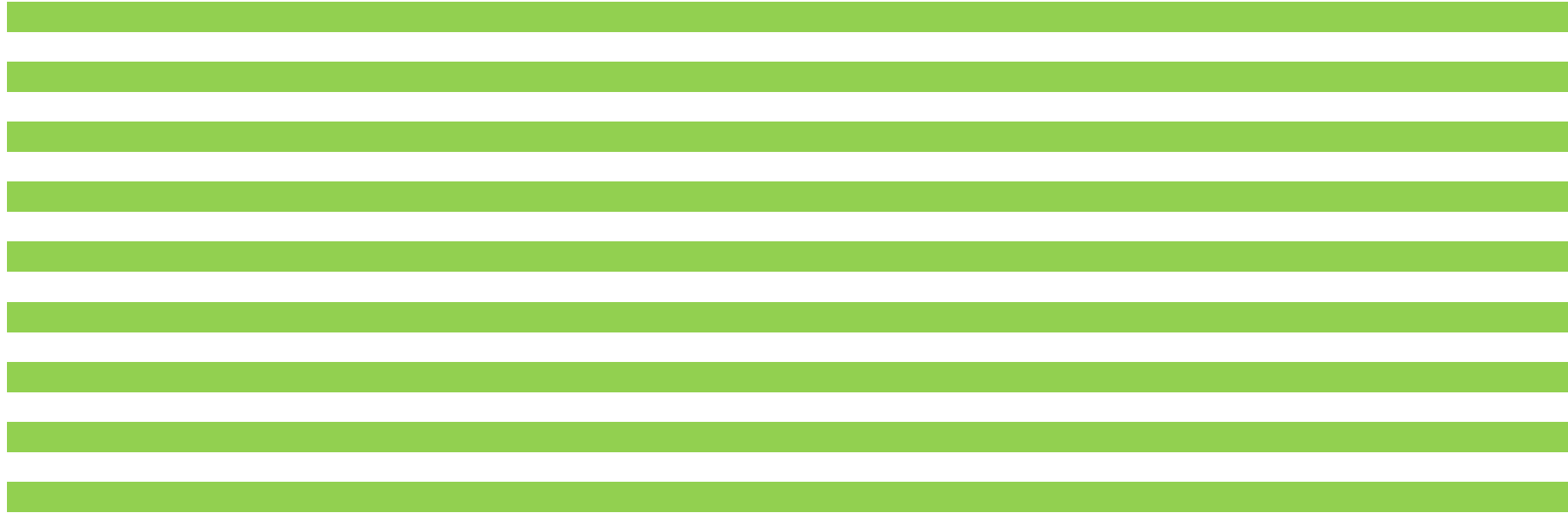
TAB]



TAB]



TAB]



TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

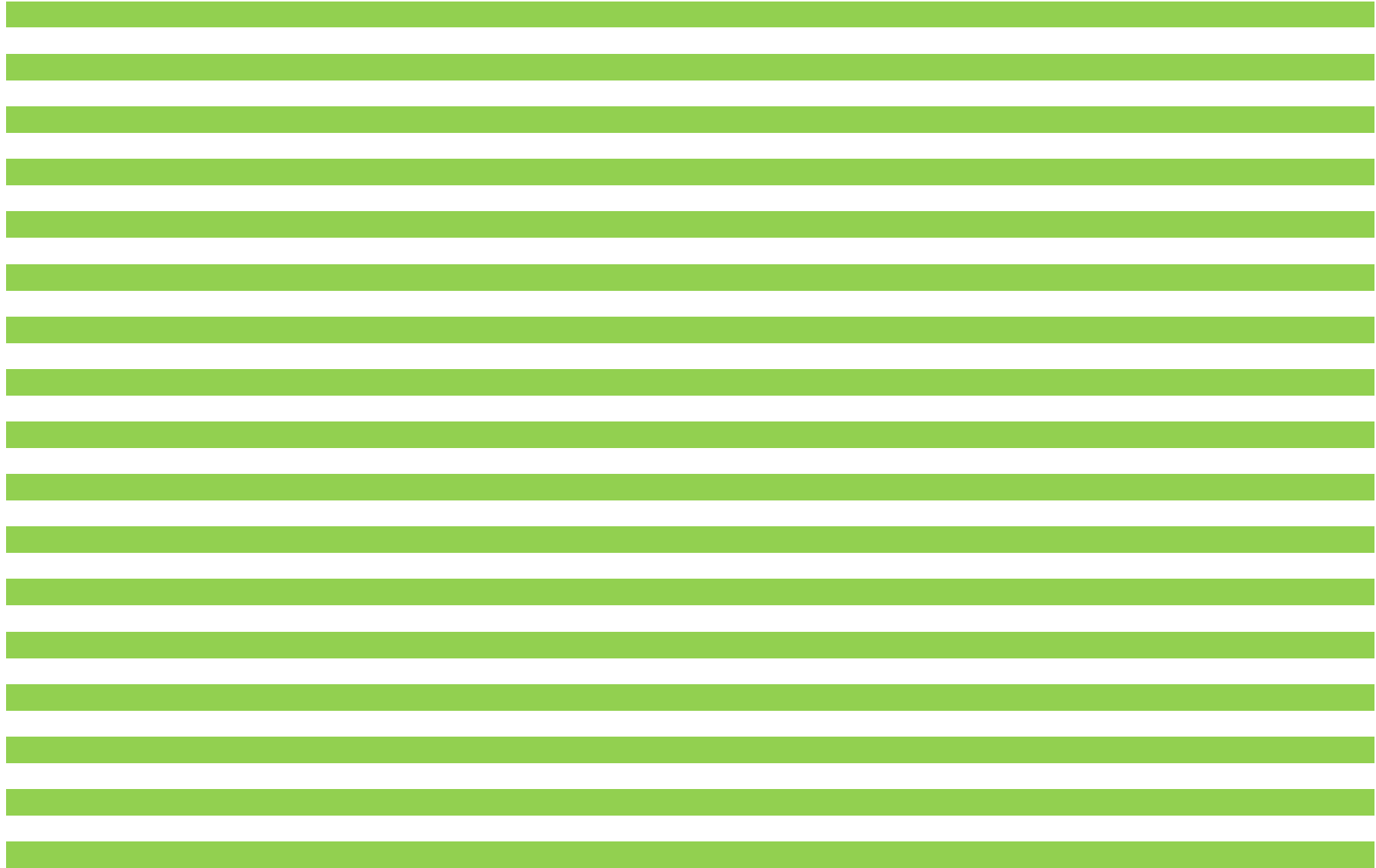
TAB]

TAB]

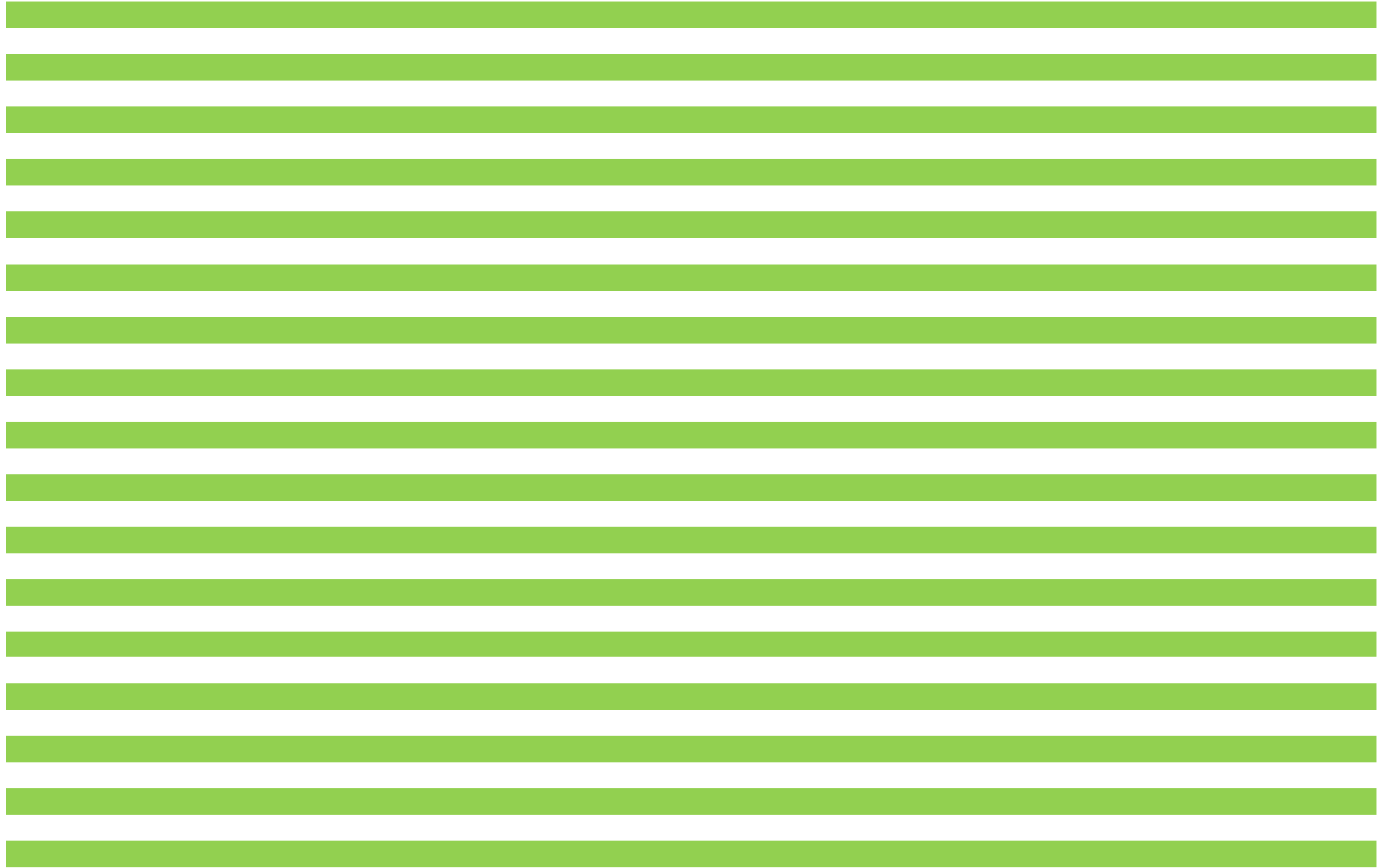
TAB]

TAB]

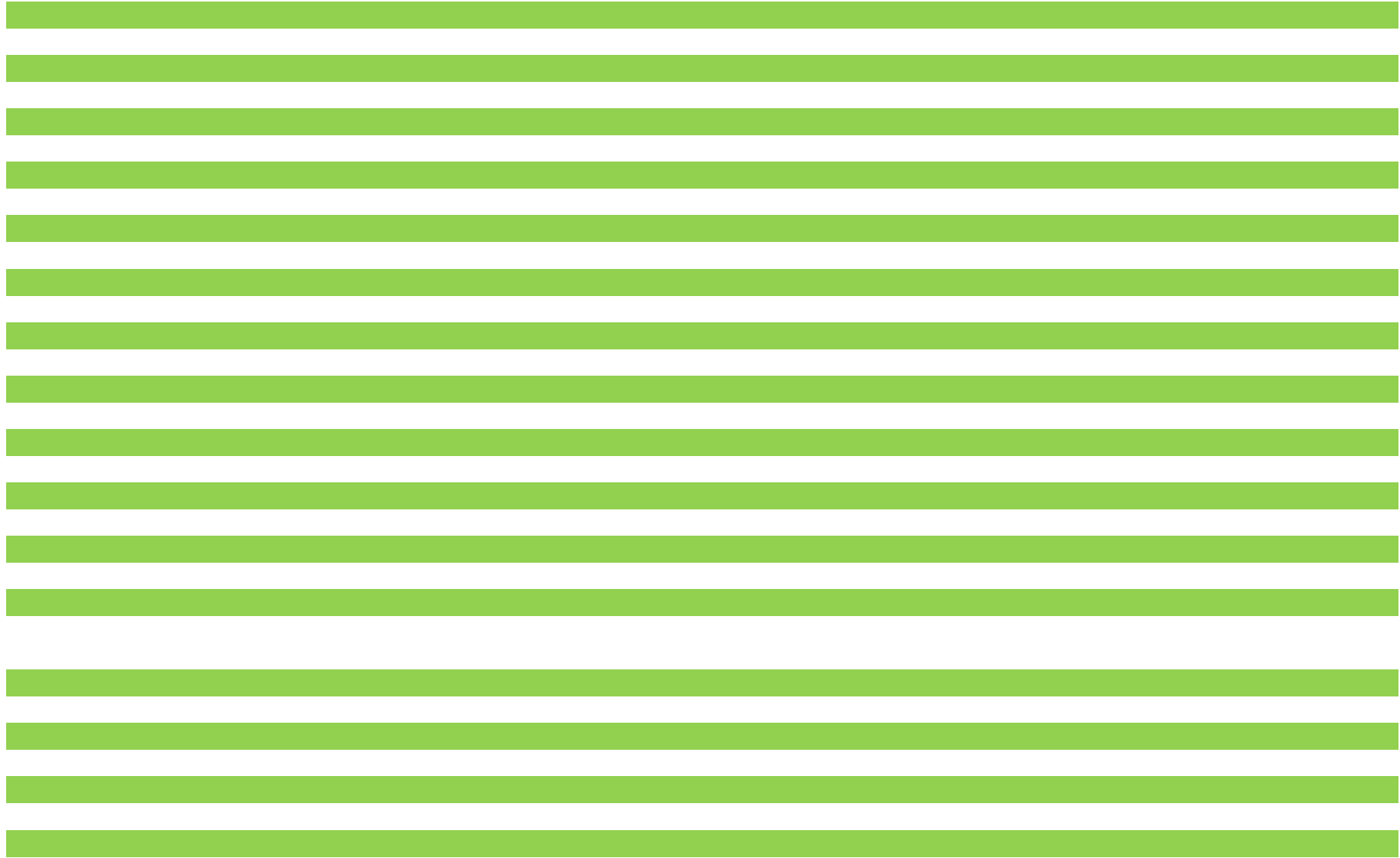
TAB]



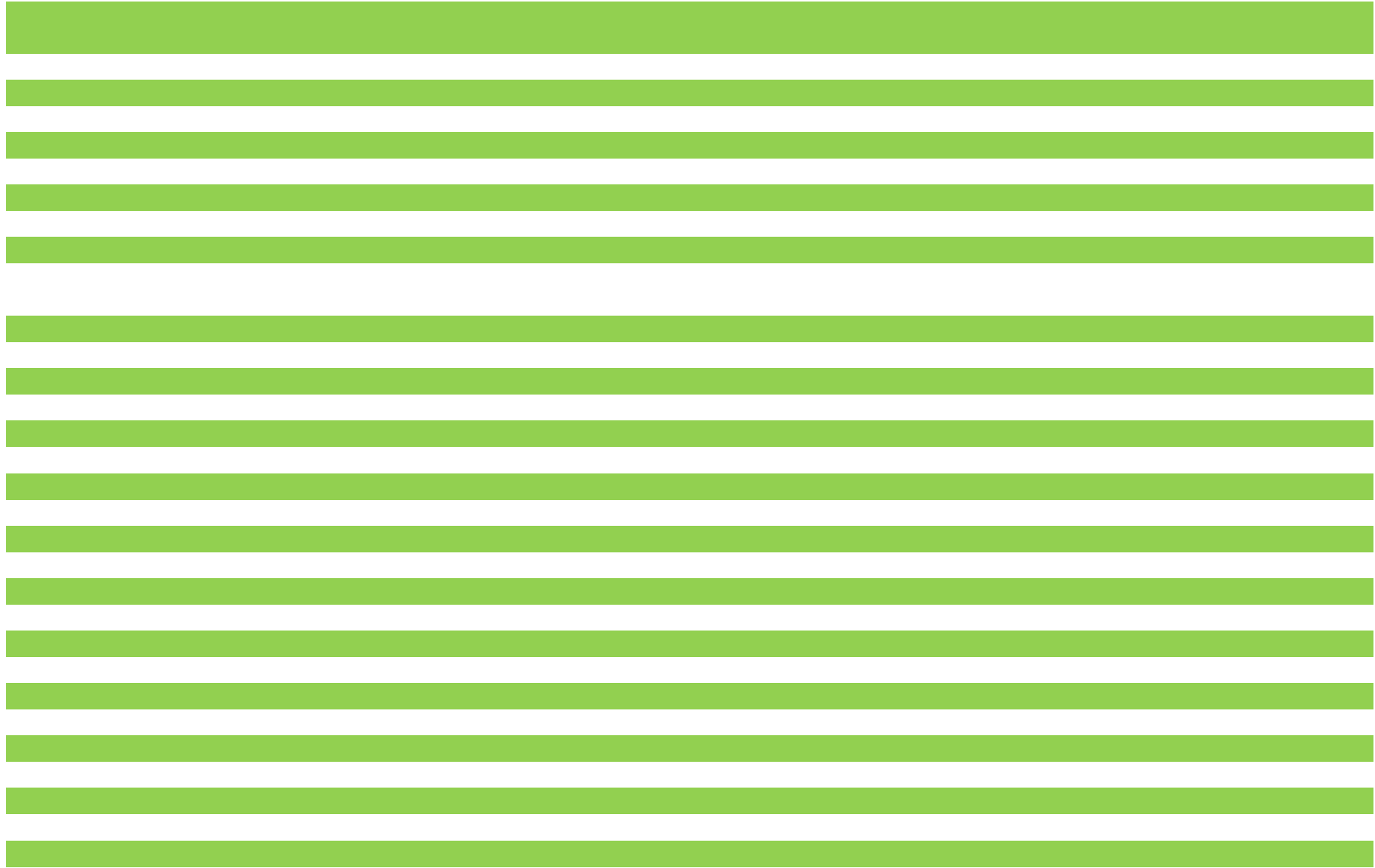
TAB]



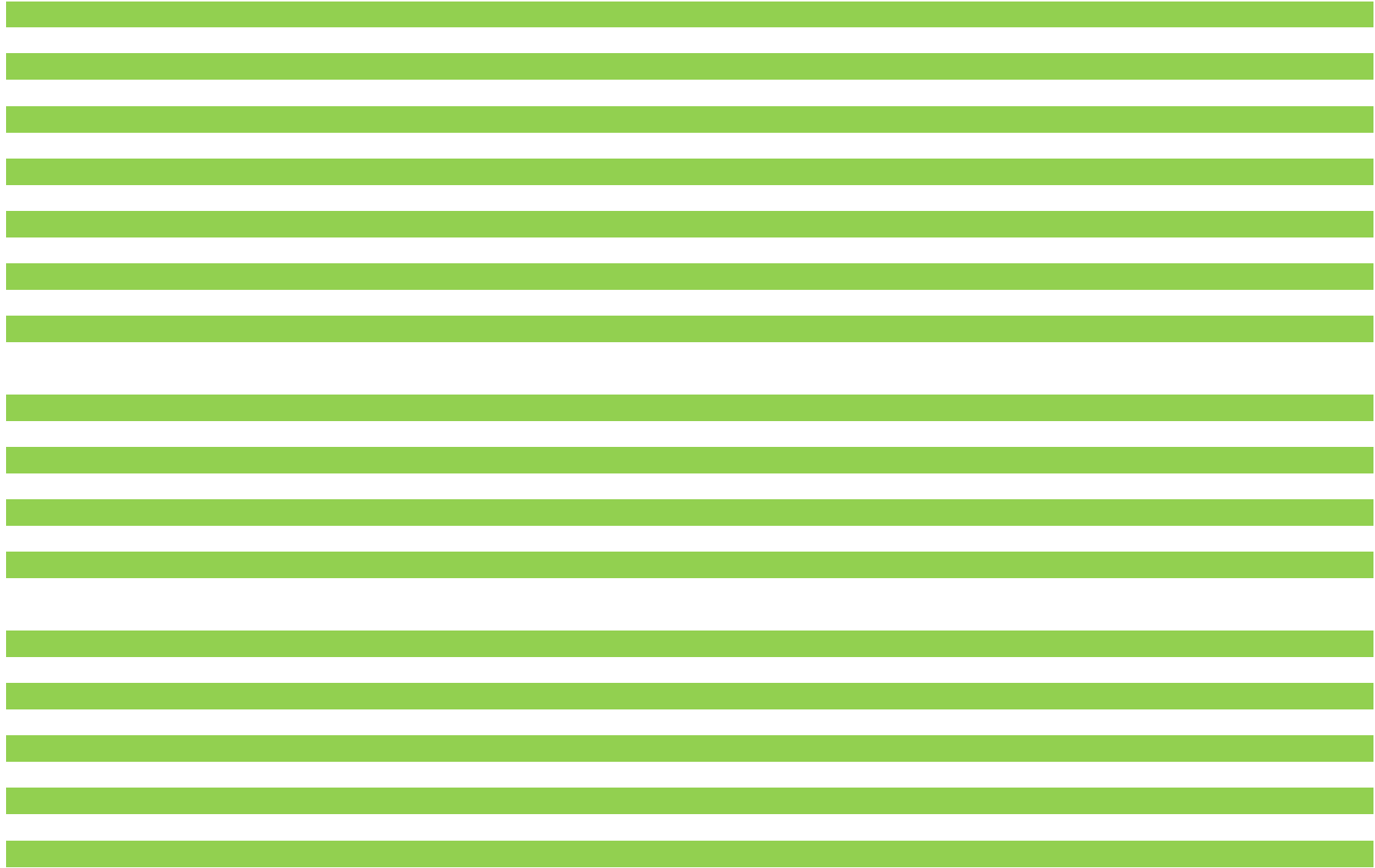
TAB]



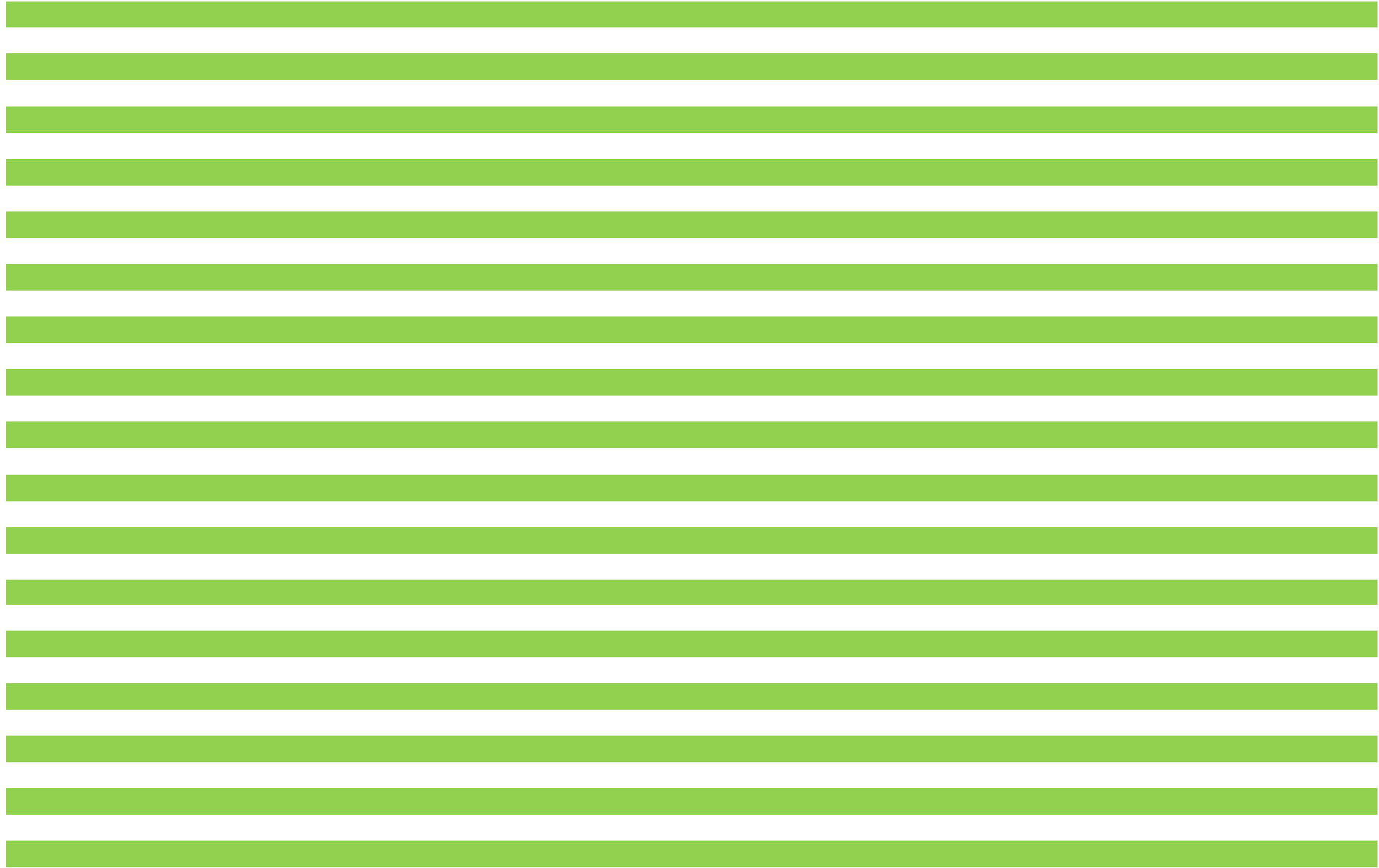
TAB]



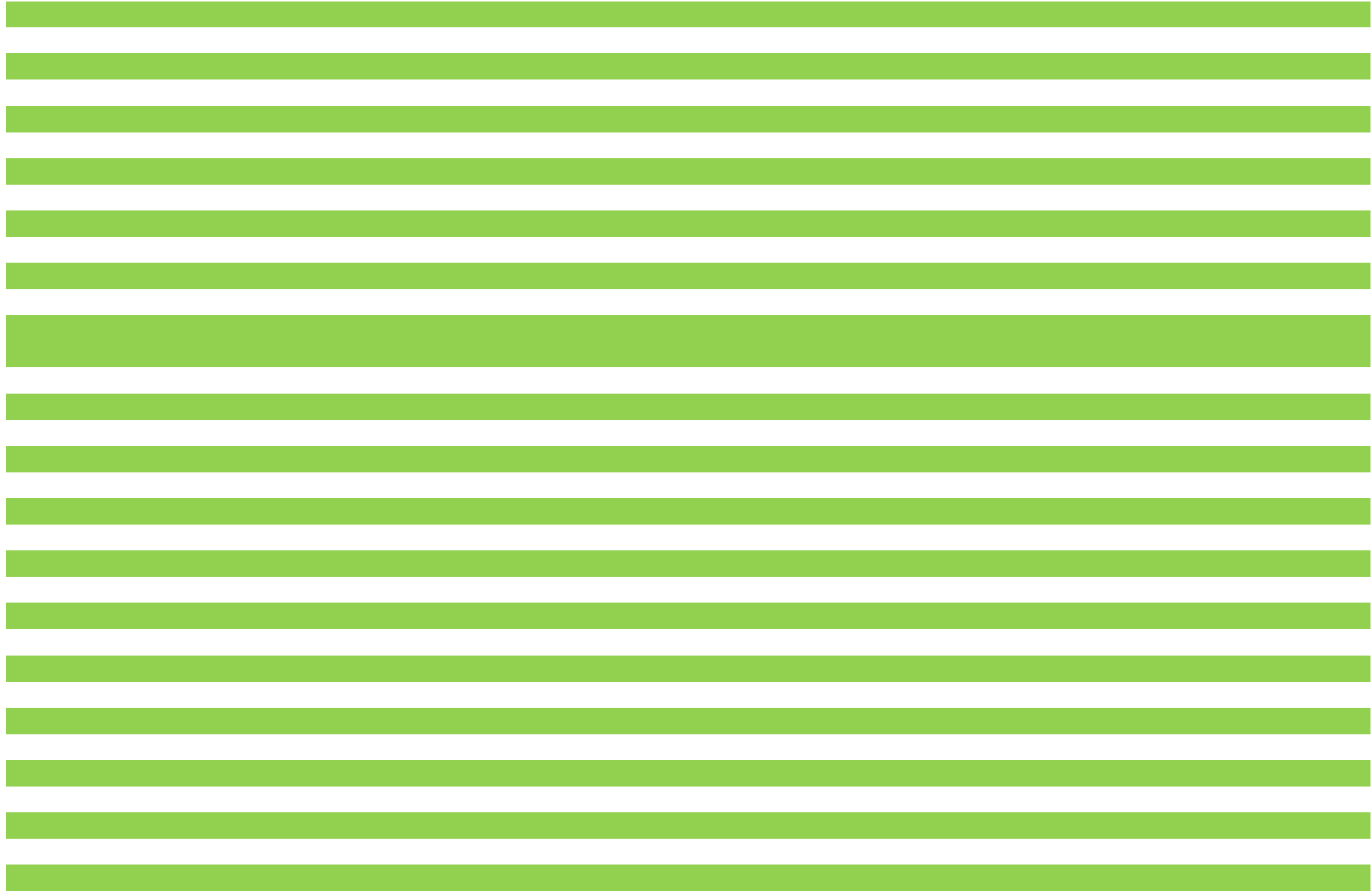
TAB]



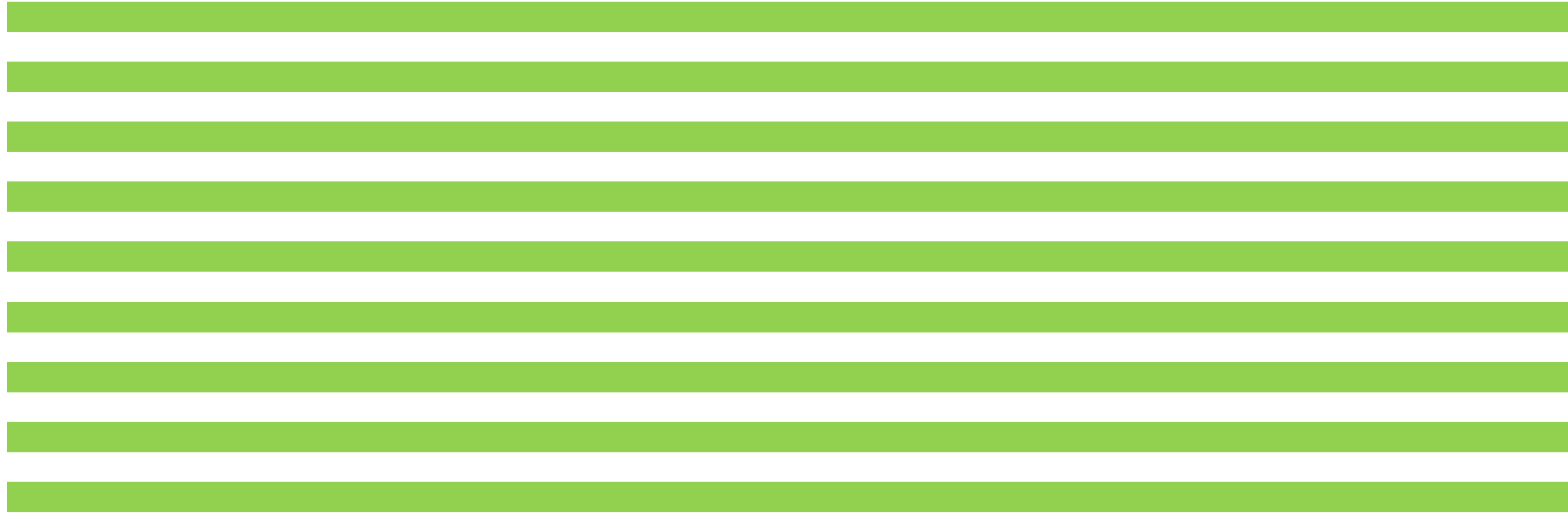
TAB]



TAB]



TAB]



TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

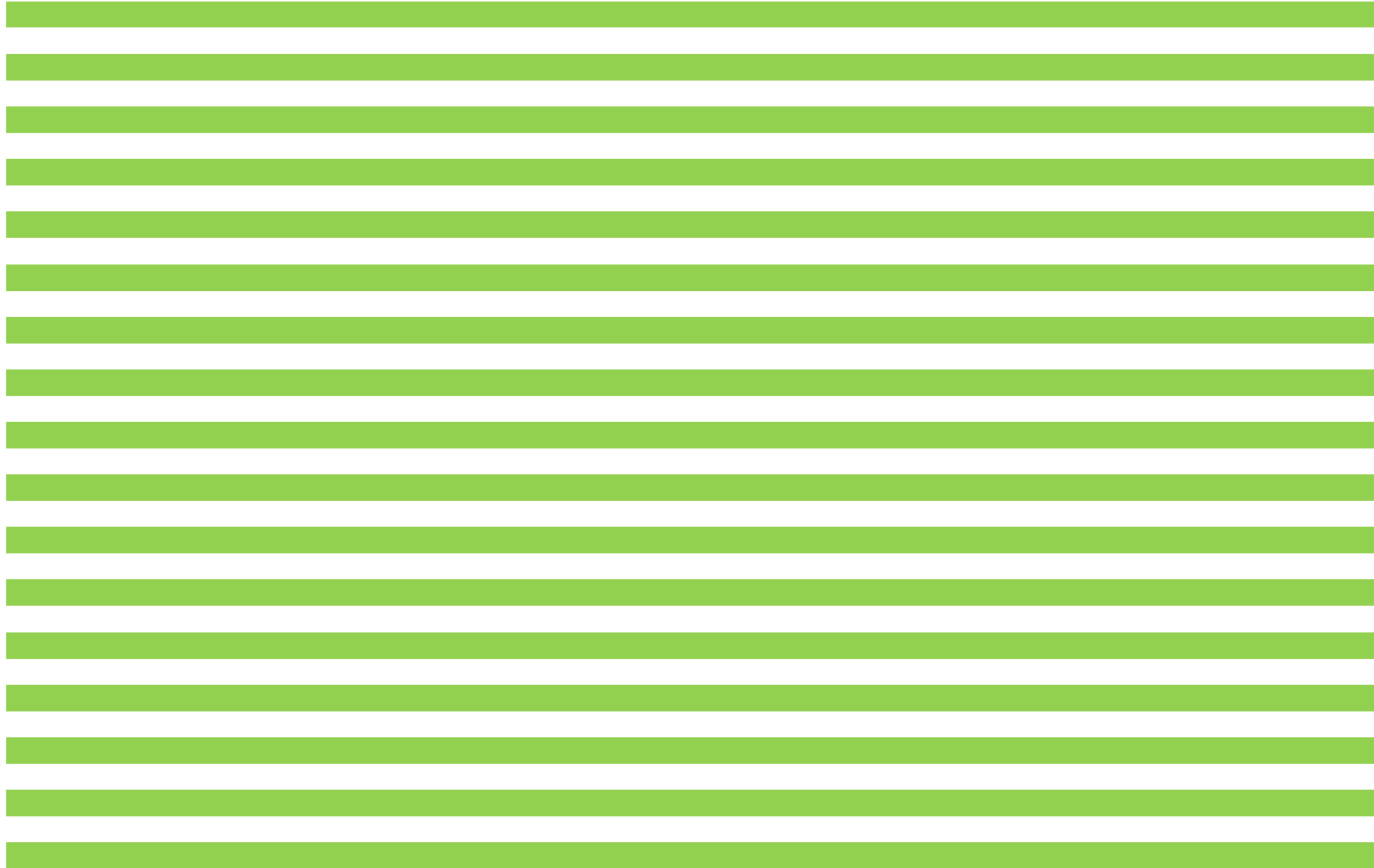
TAB]

TAB]

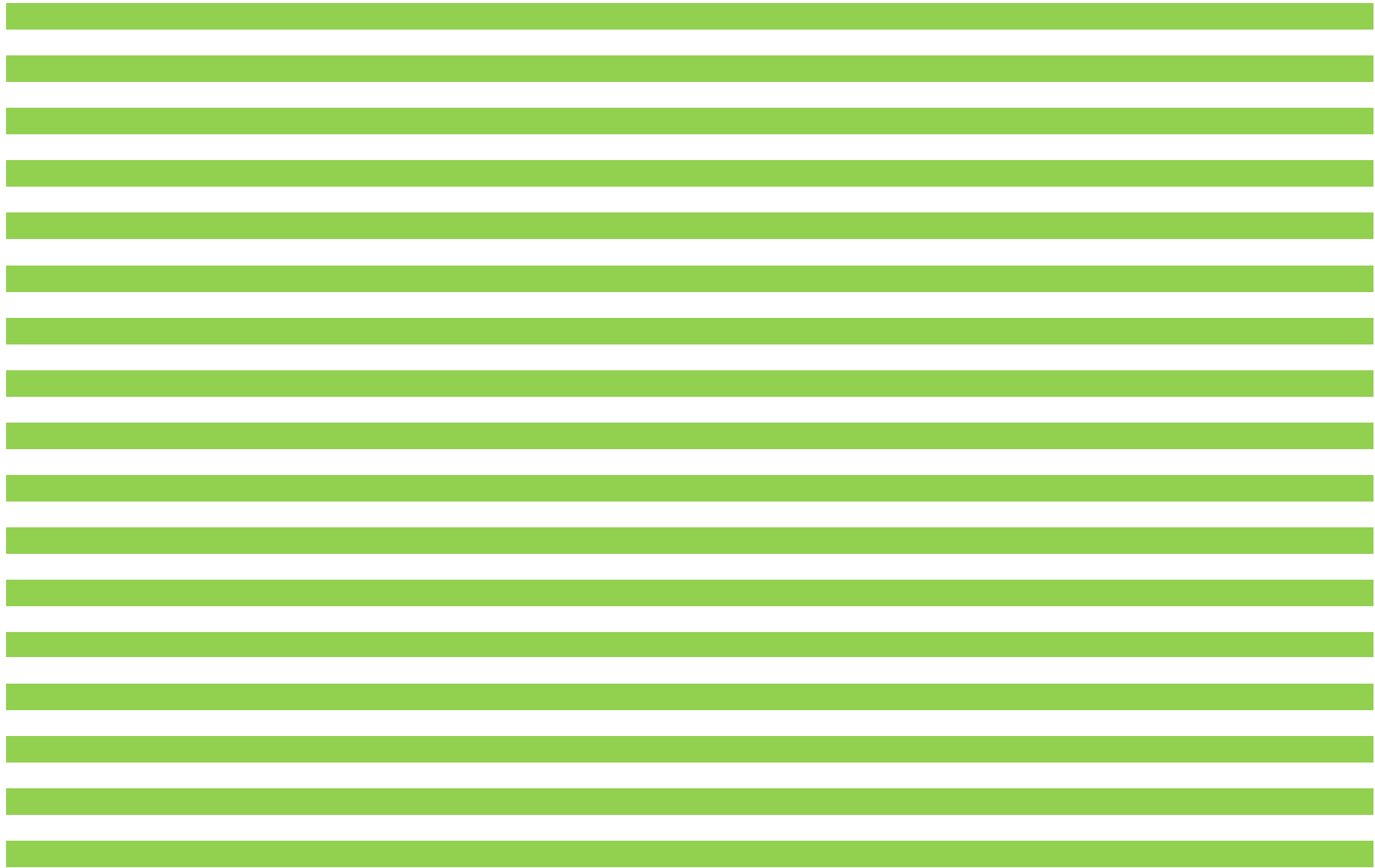
TAB]

TAB]

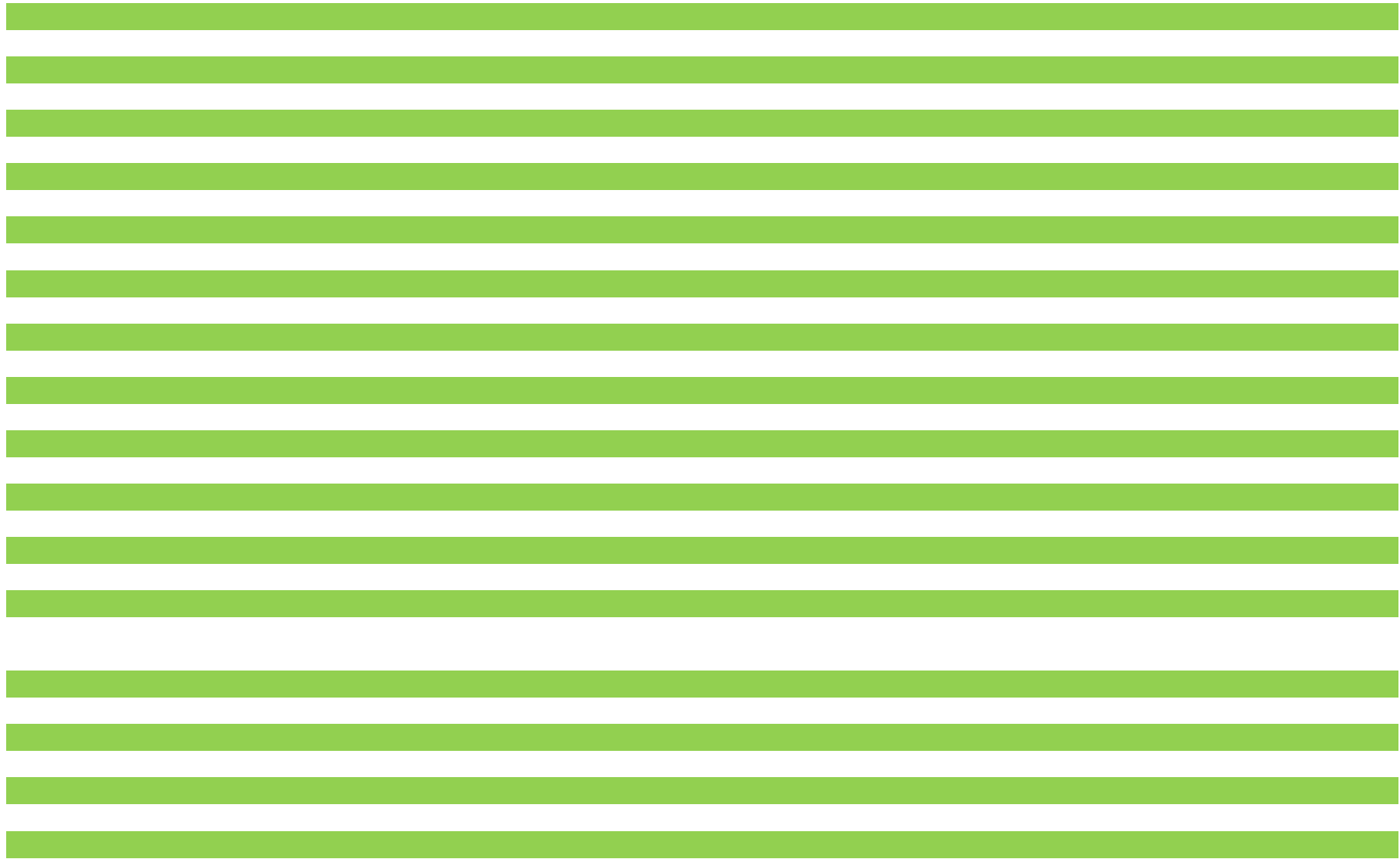
TAB]



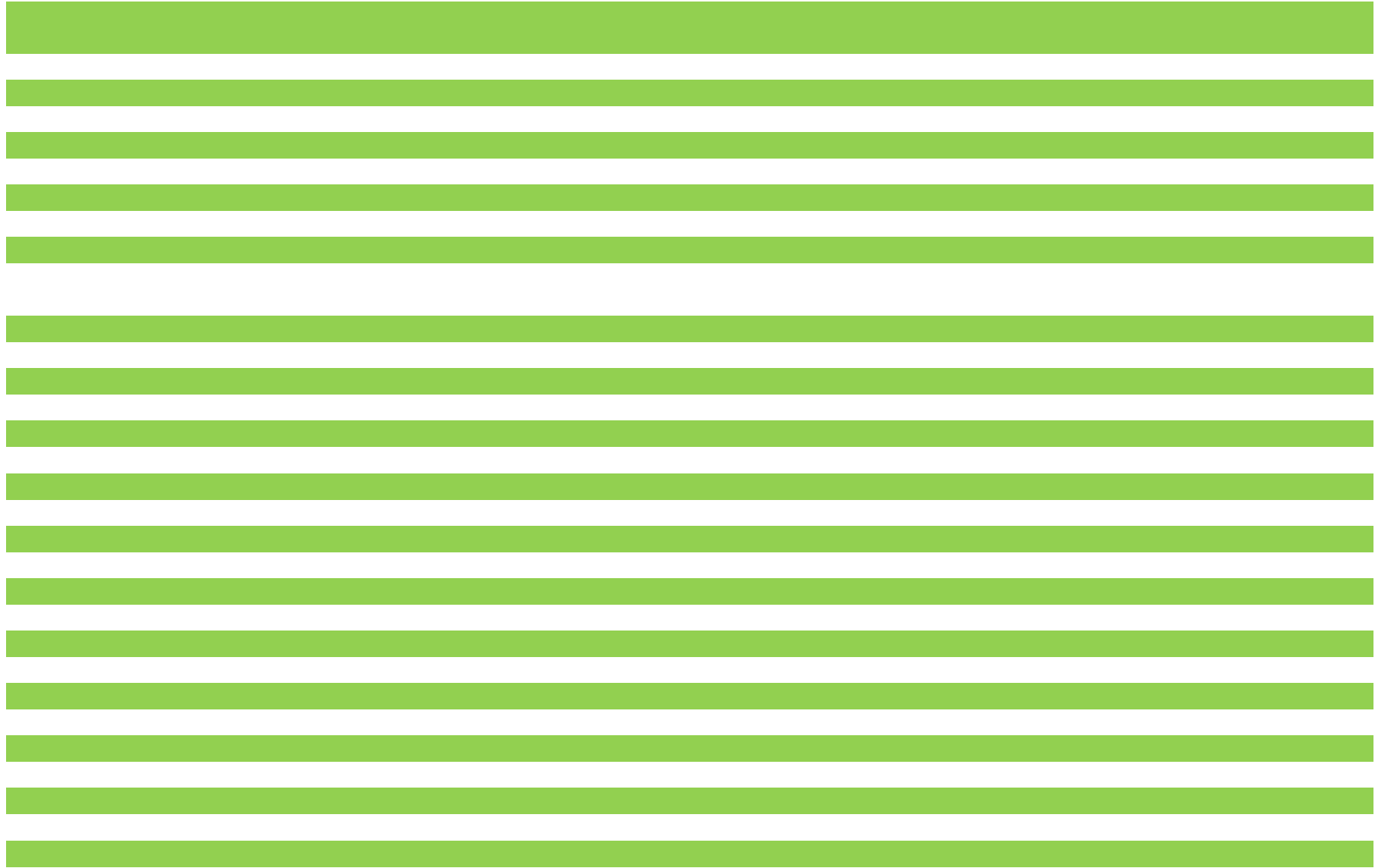
TAB]



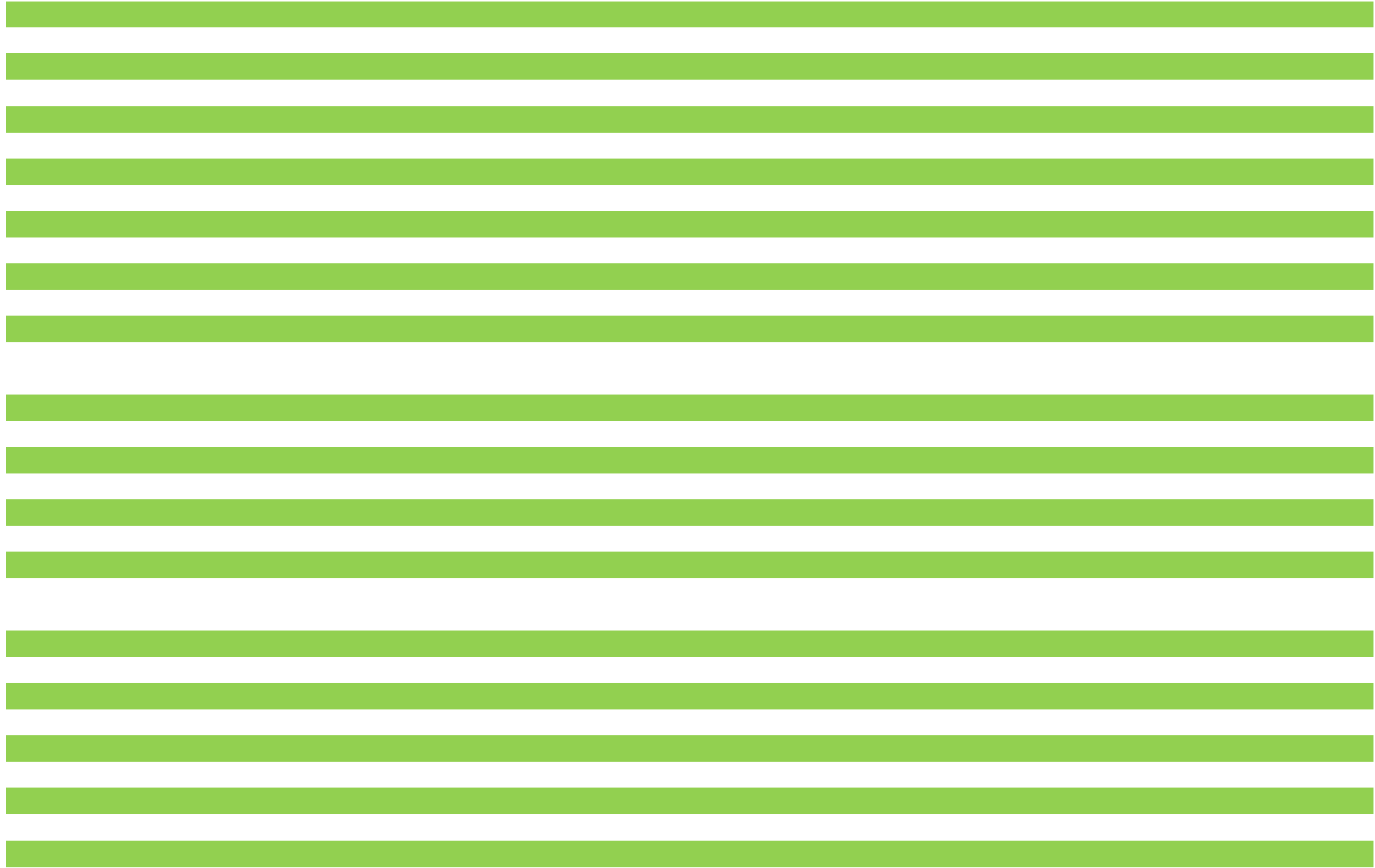
TAB]



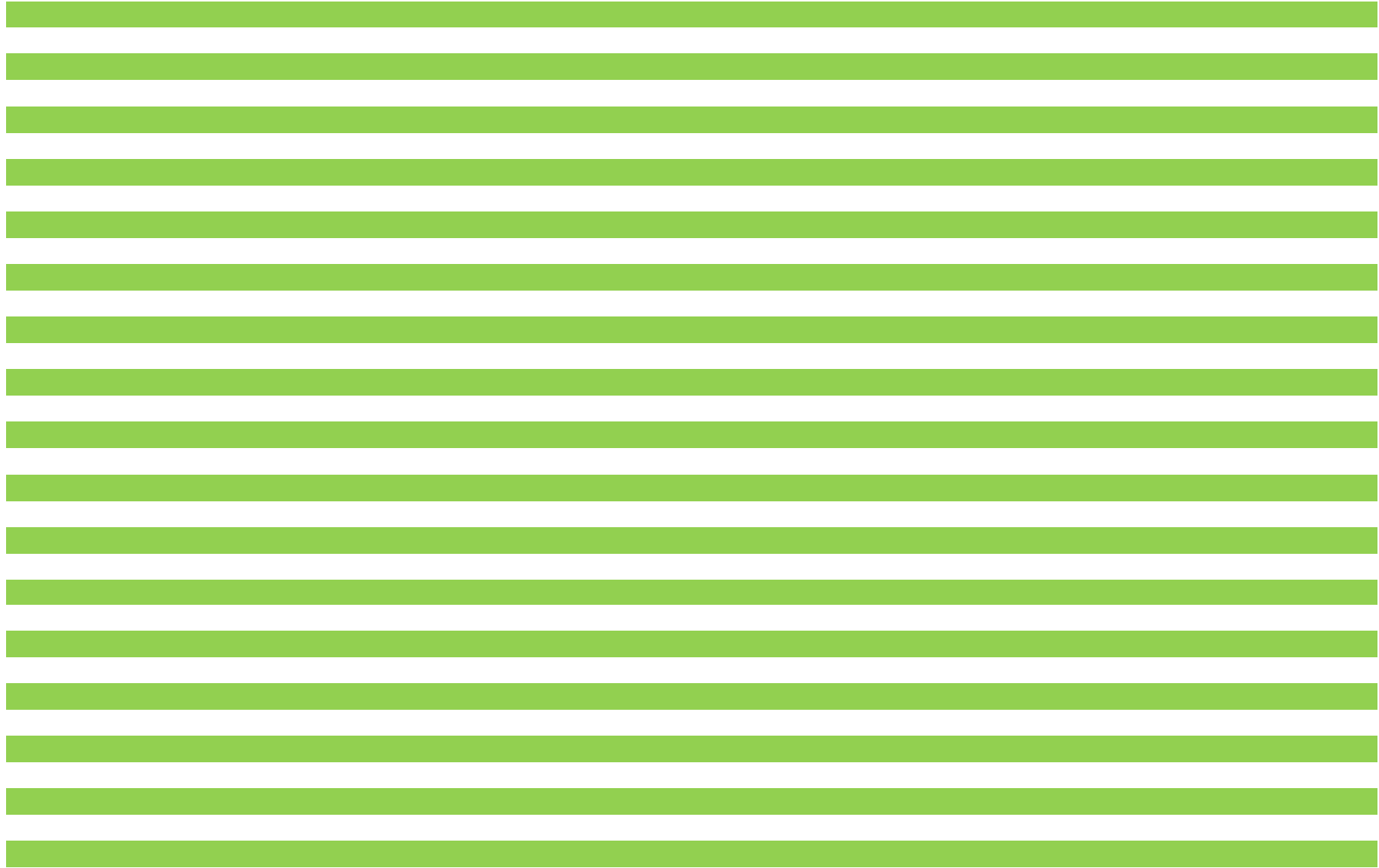
TAB]



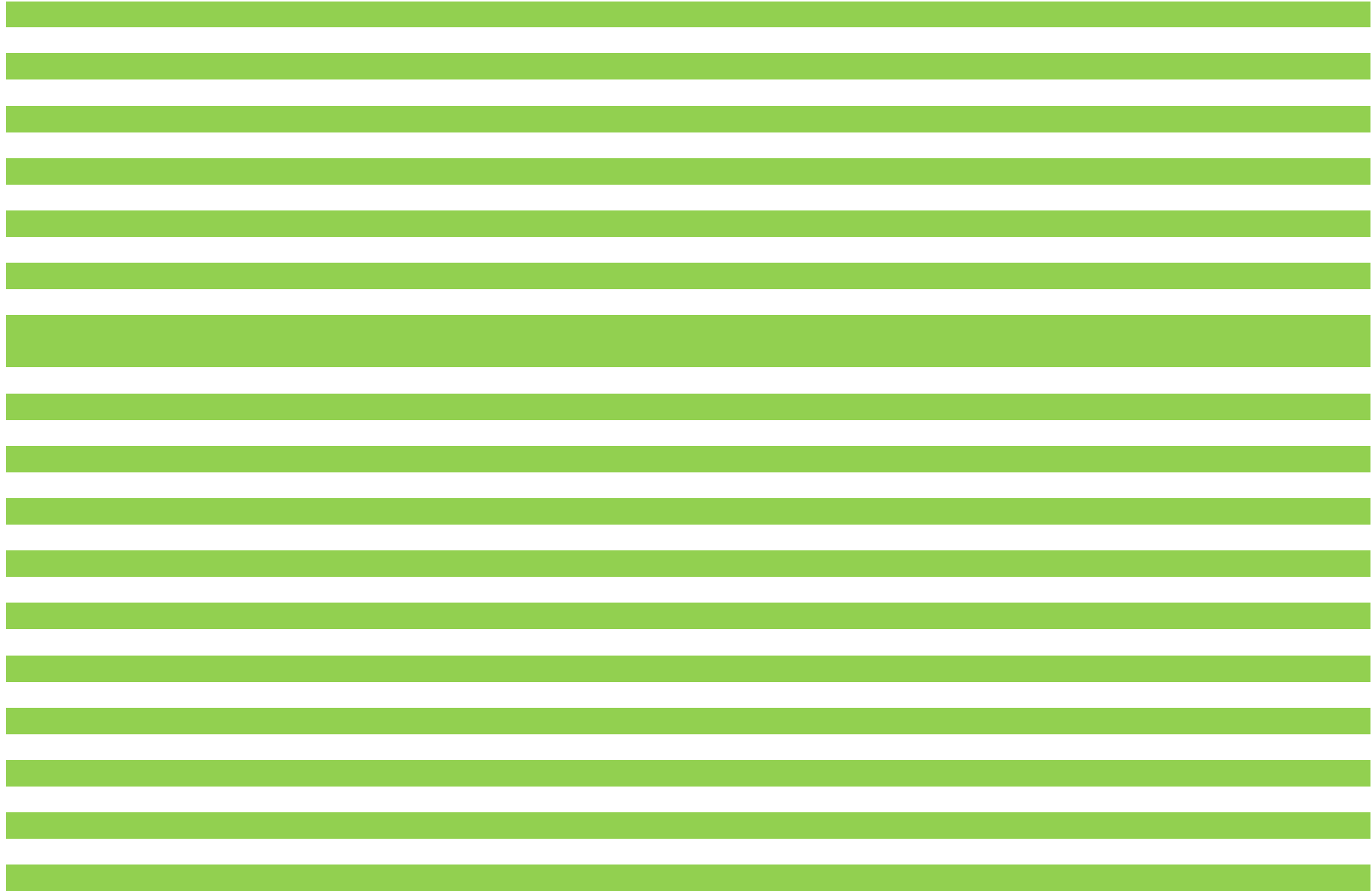
TAB]



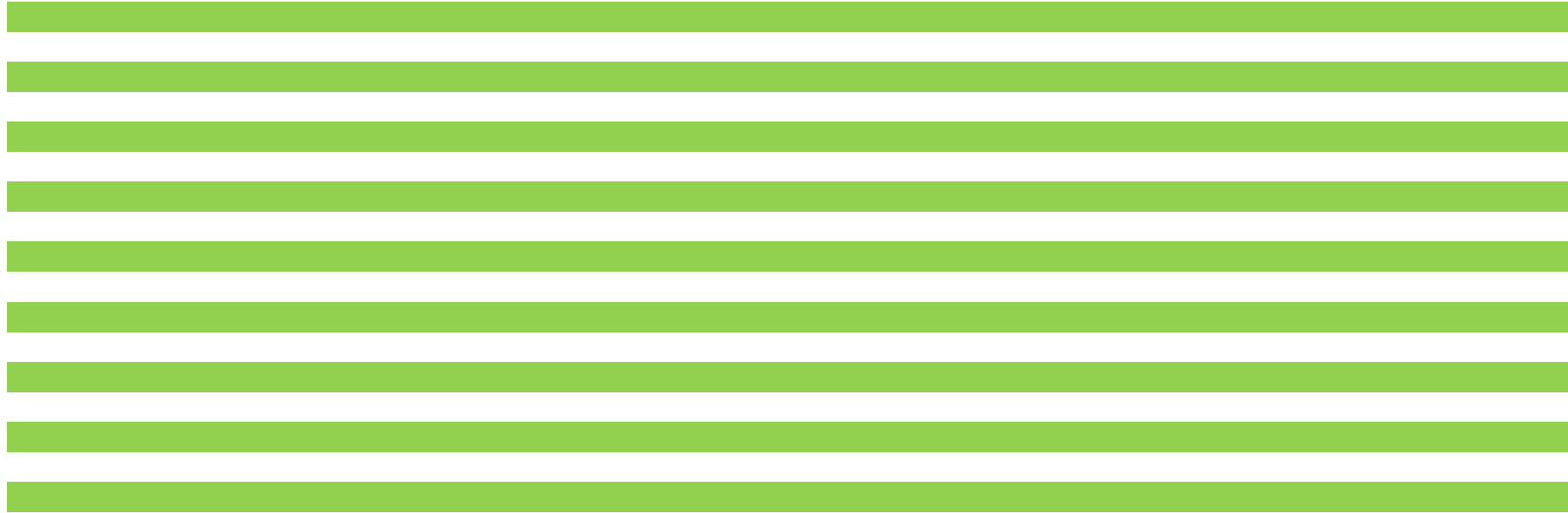
TAB]



TAB]



TAB]



TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

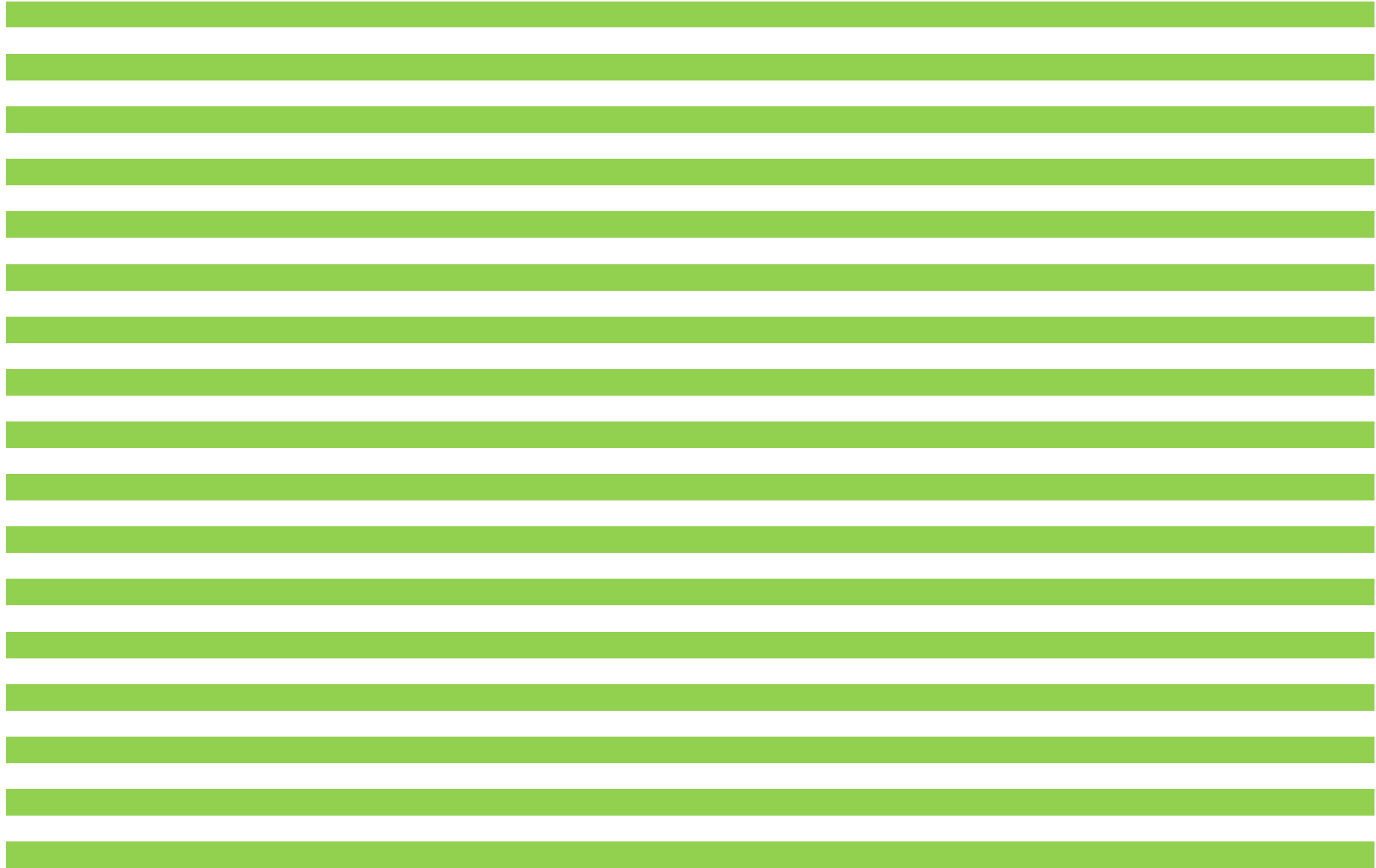
TAB]

TAB]

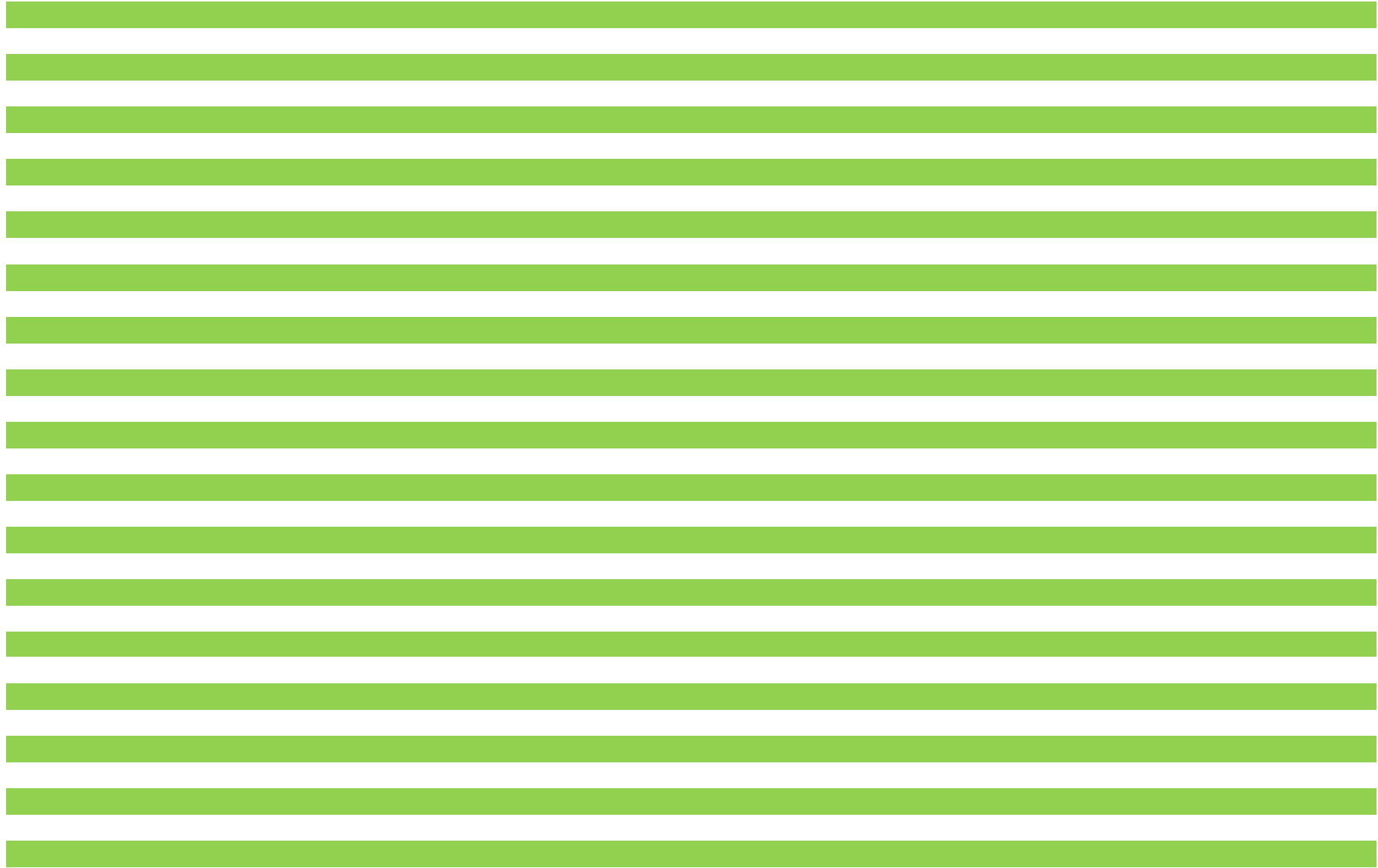
TAB]

TAB]

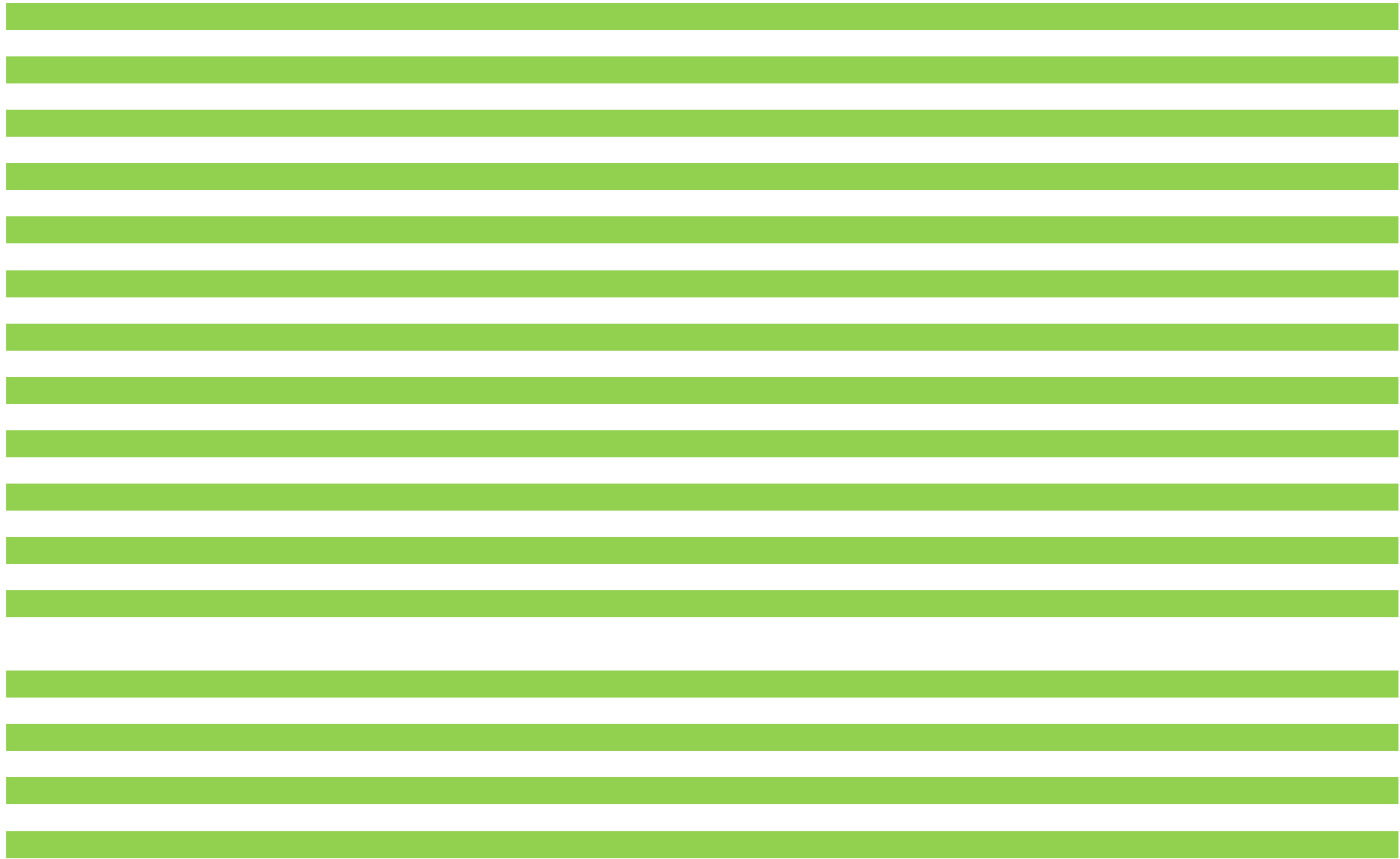
TAB]



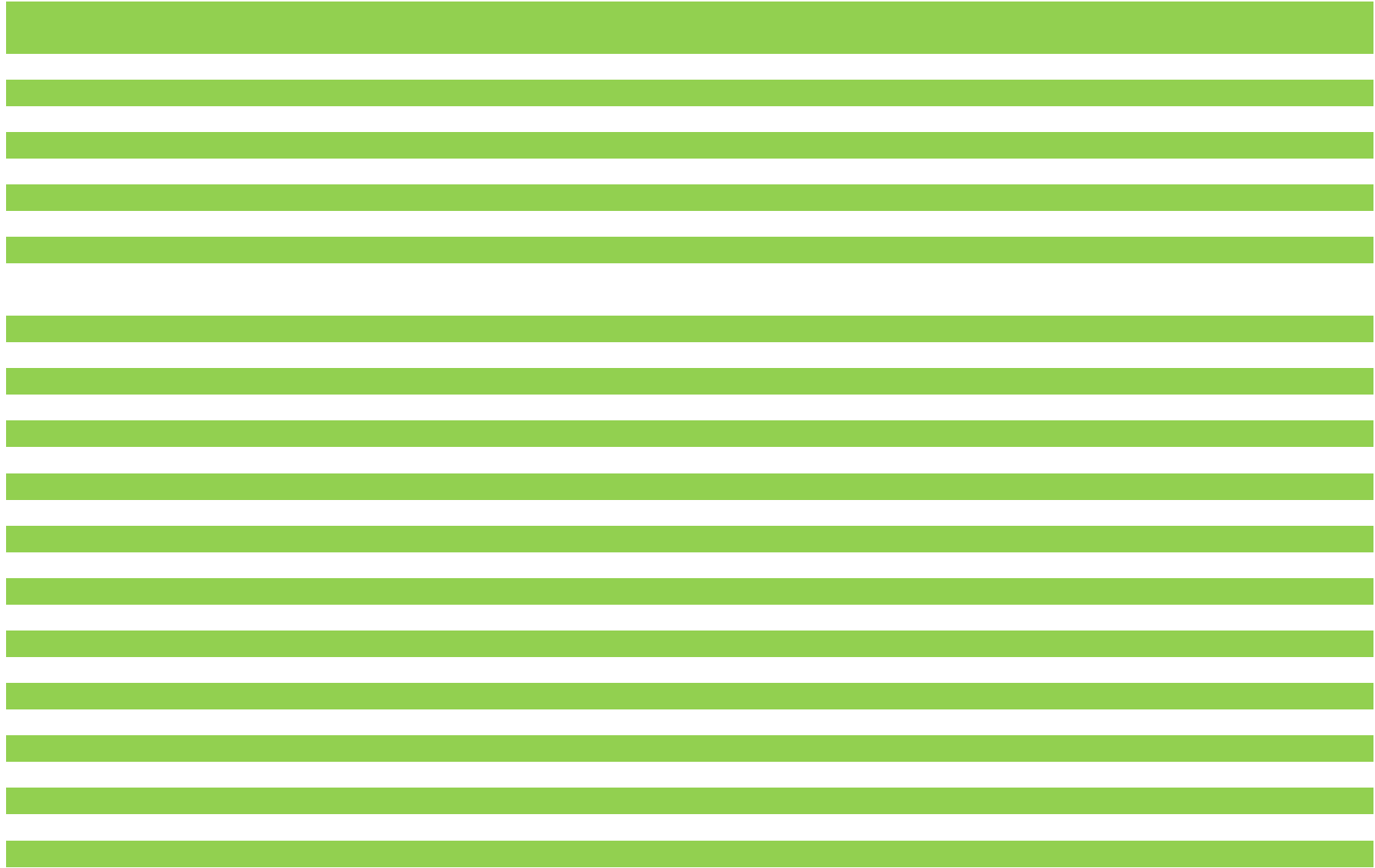
TAB]



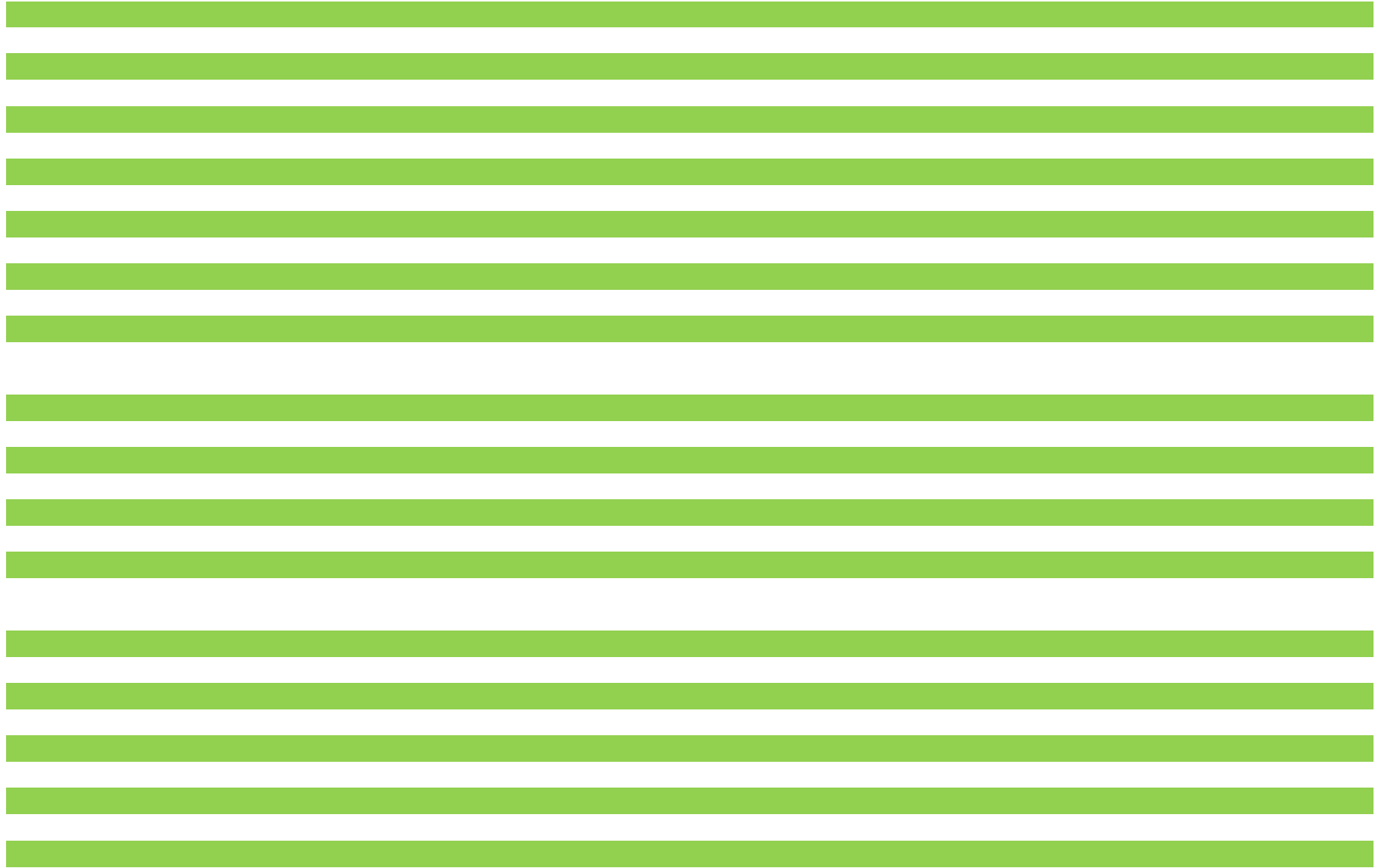
TAB]



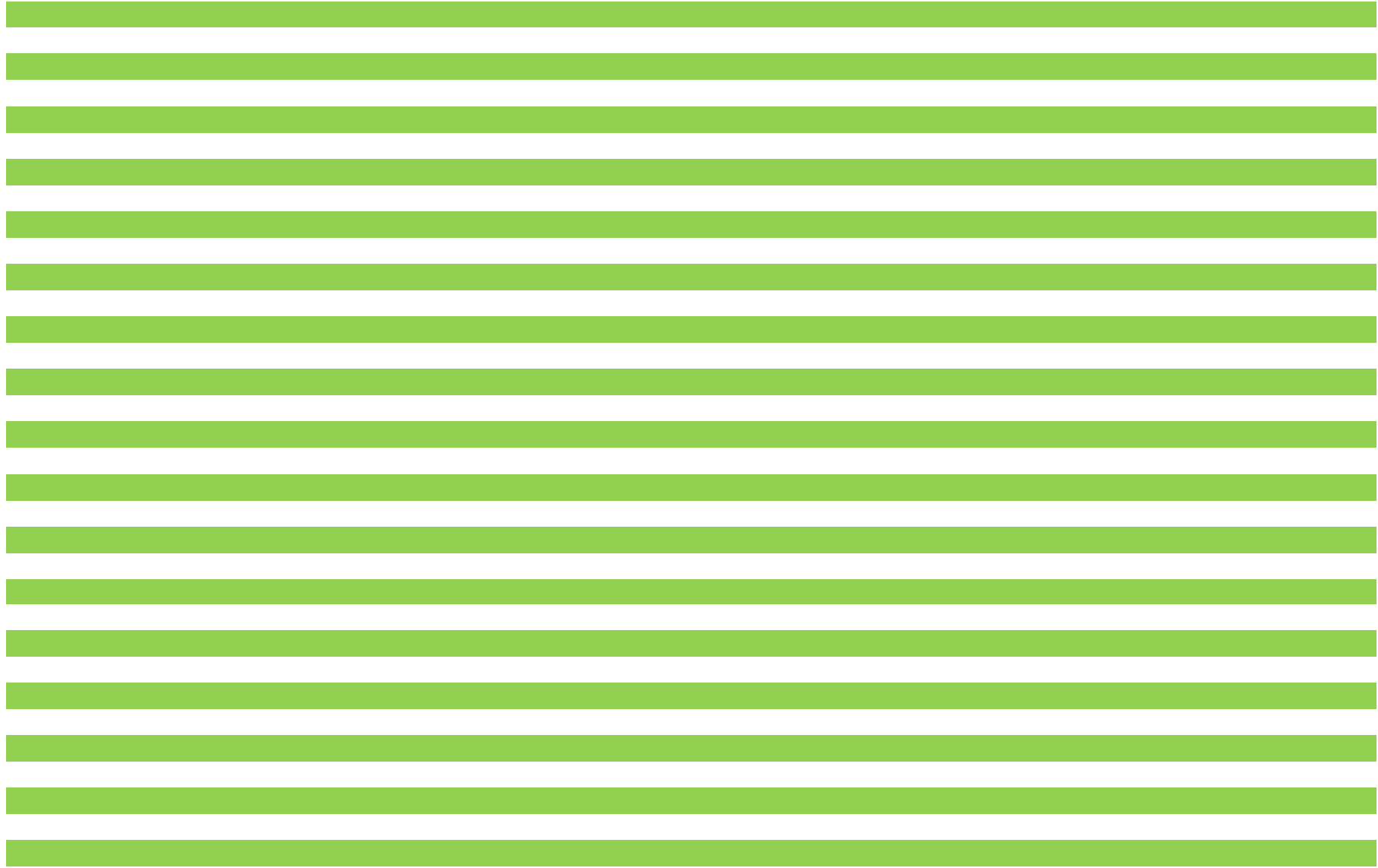
TAB]



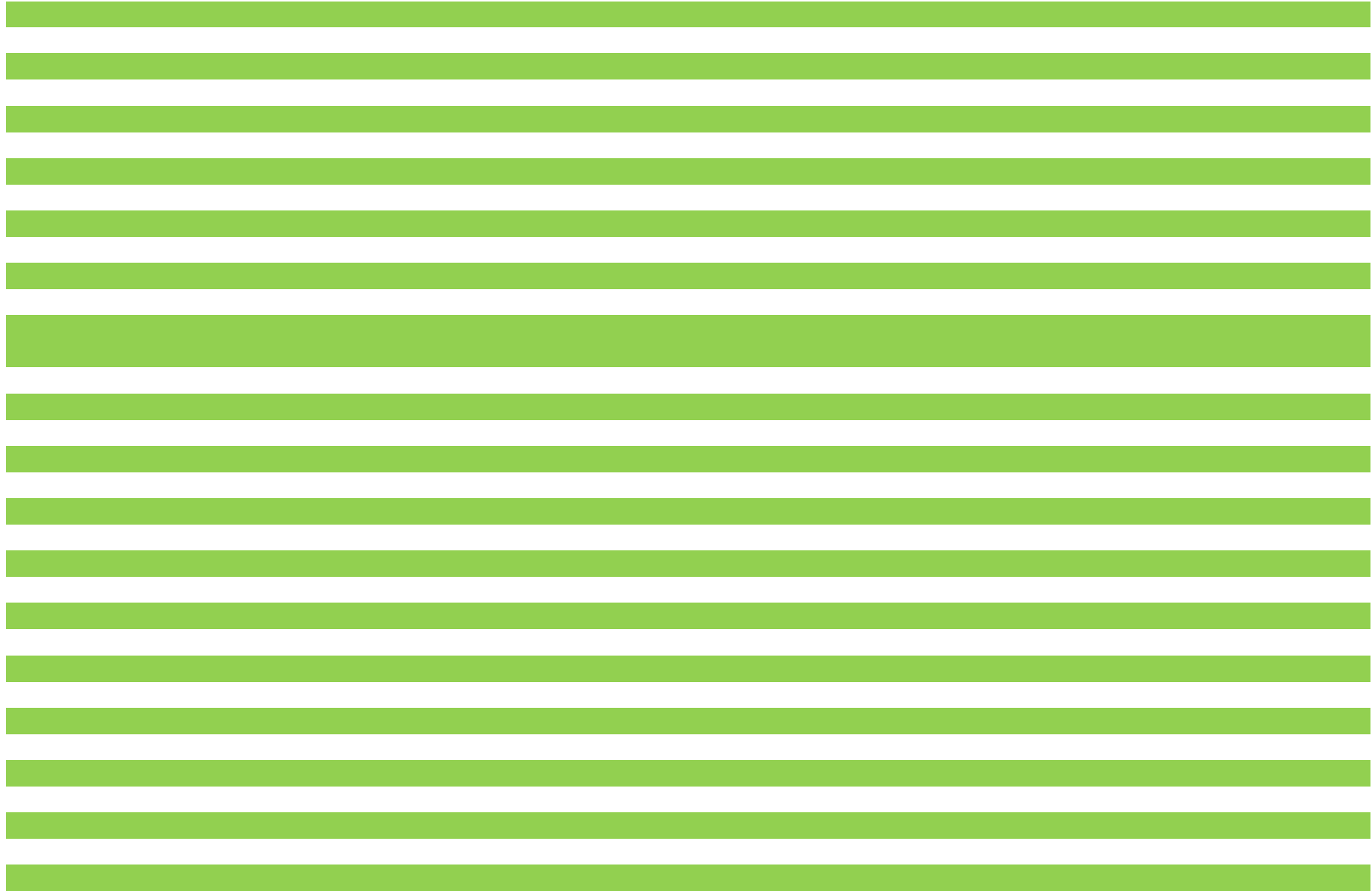
TAB]



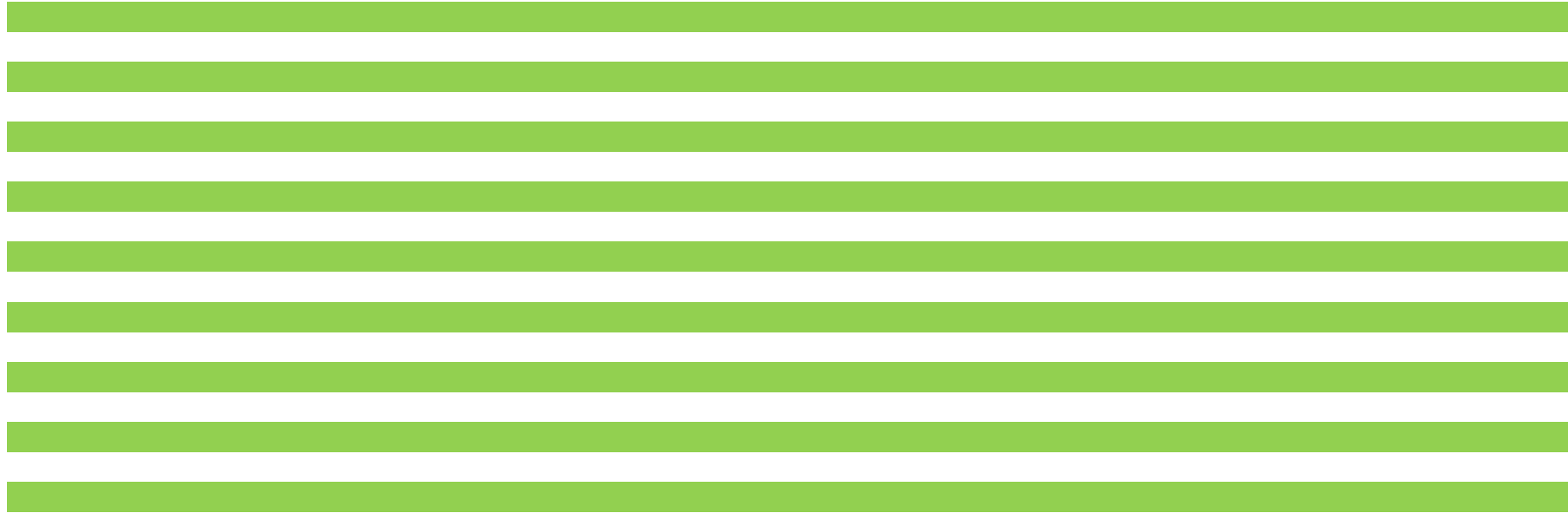
TAB]



TAB]



TAB]



TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

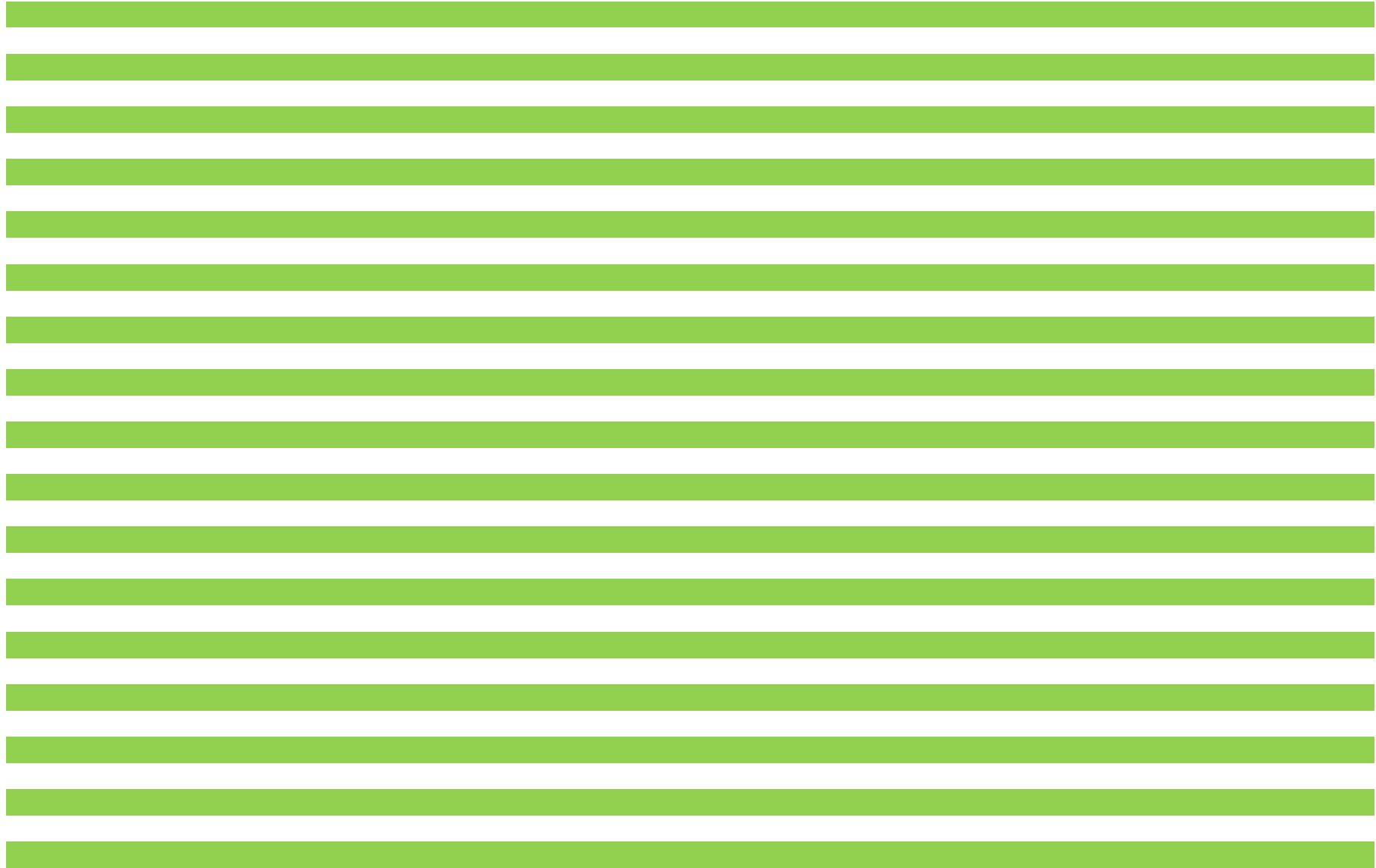
TAB]

TAB]

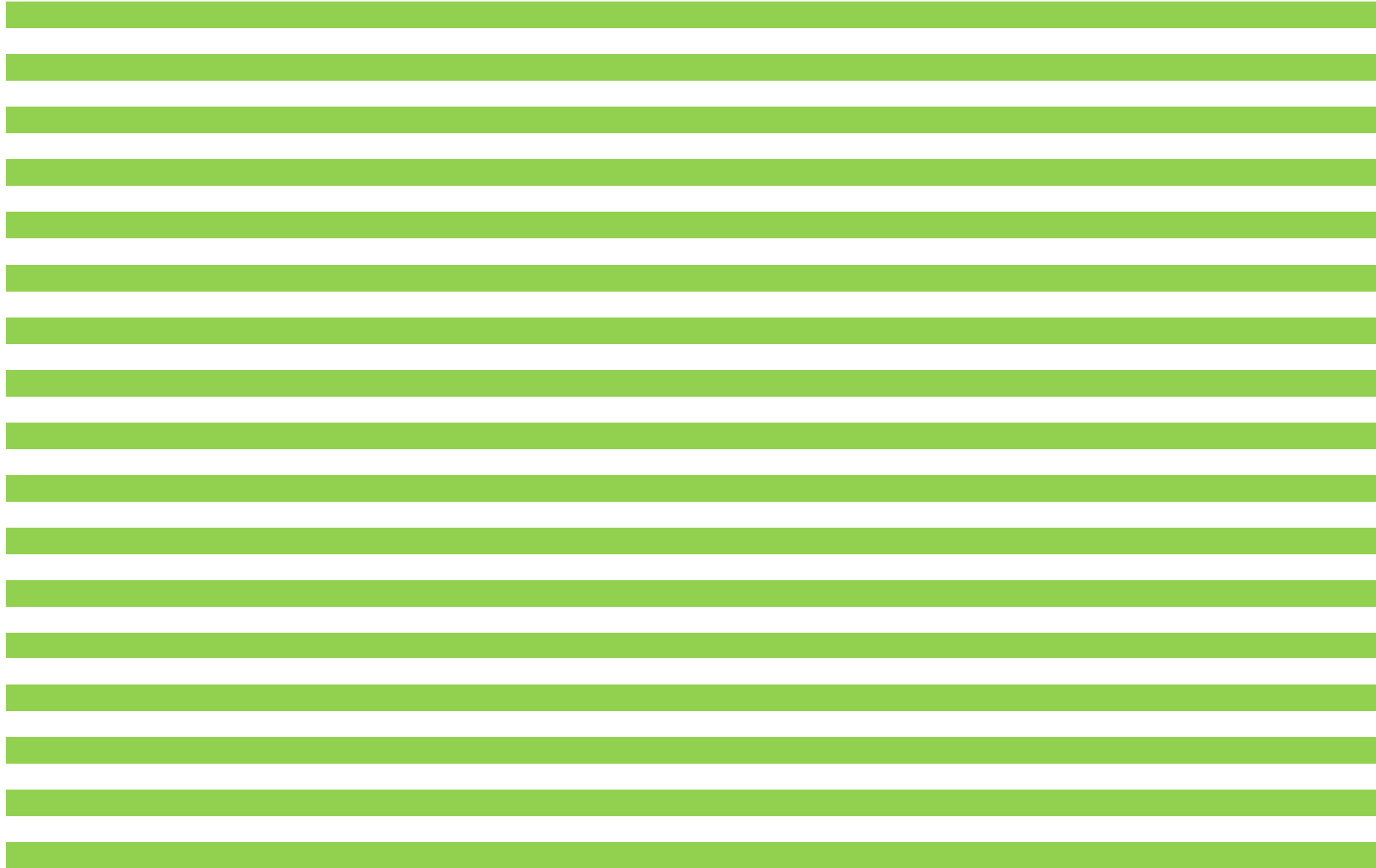
TAB]

TAB]

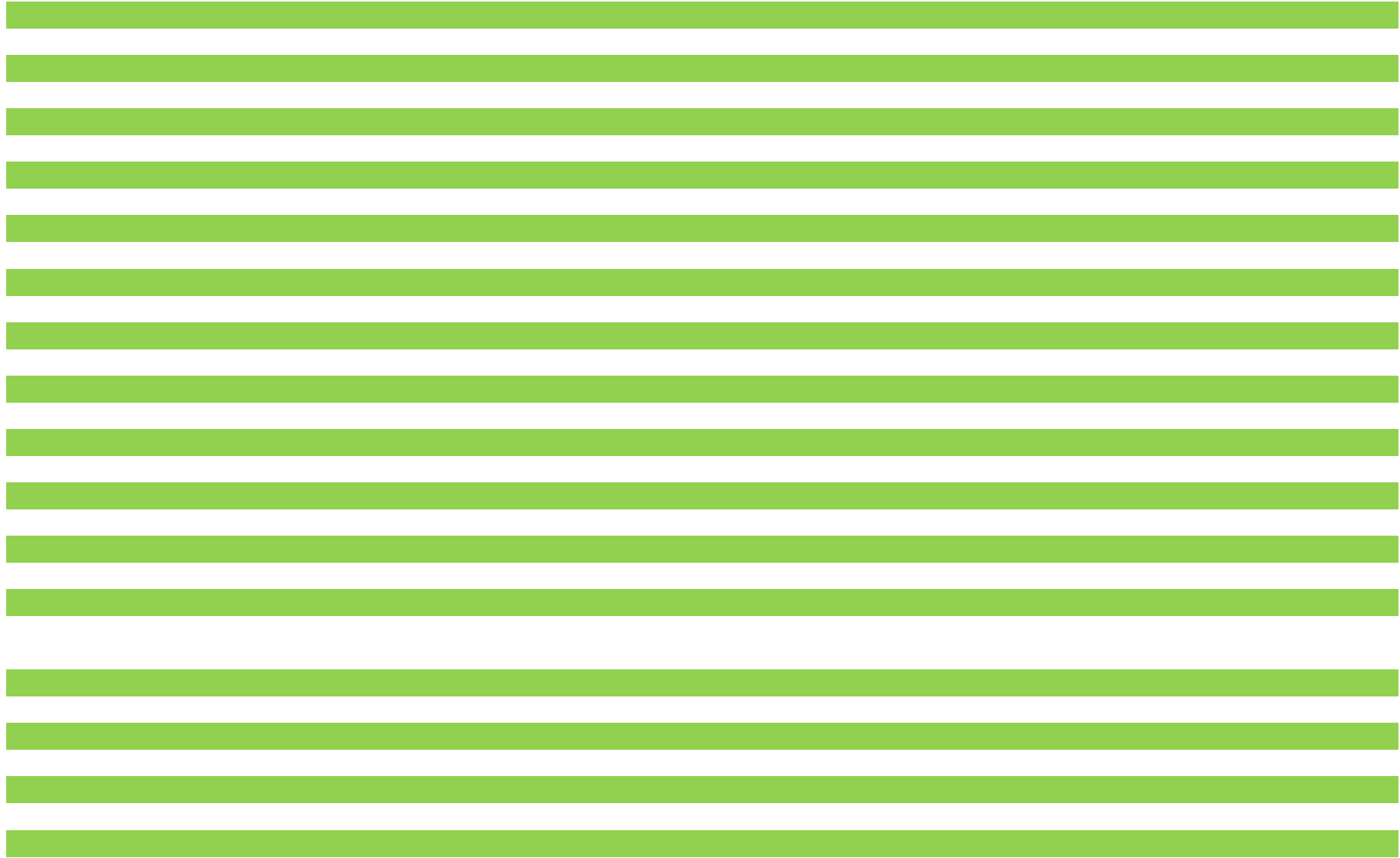
TAB]



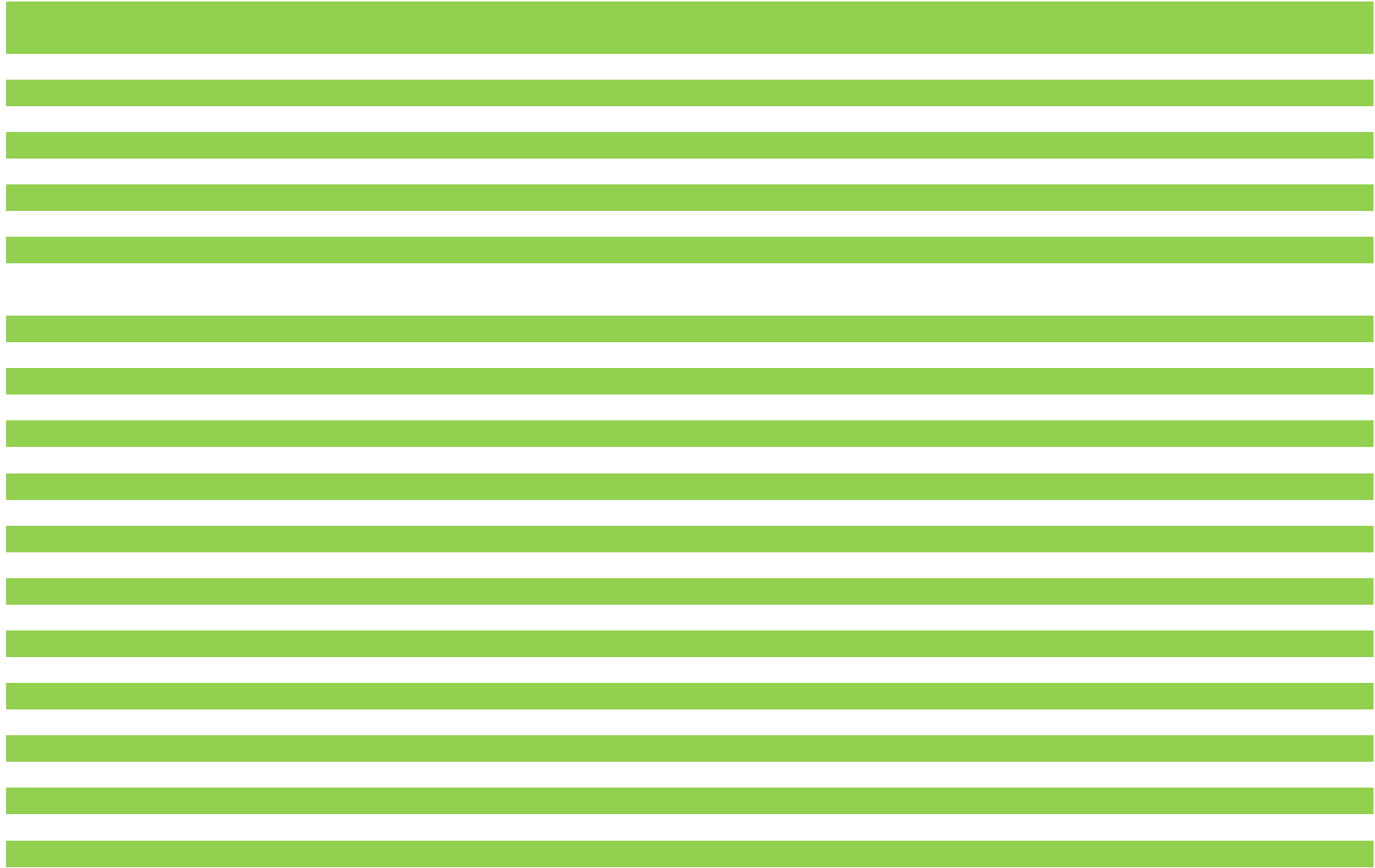
TAB]



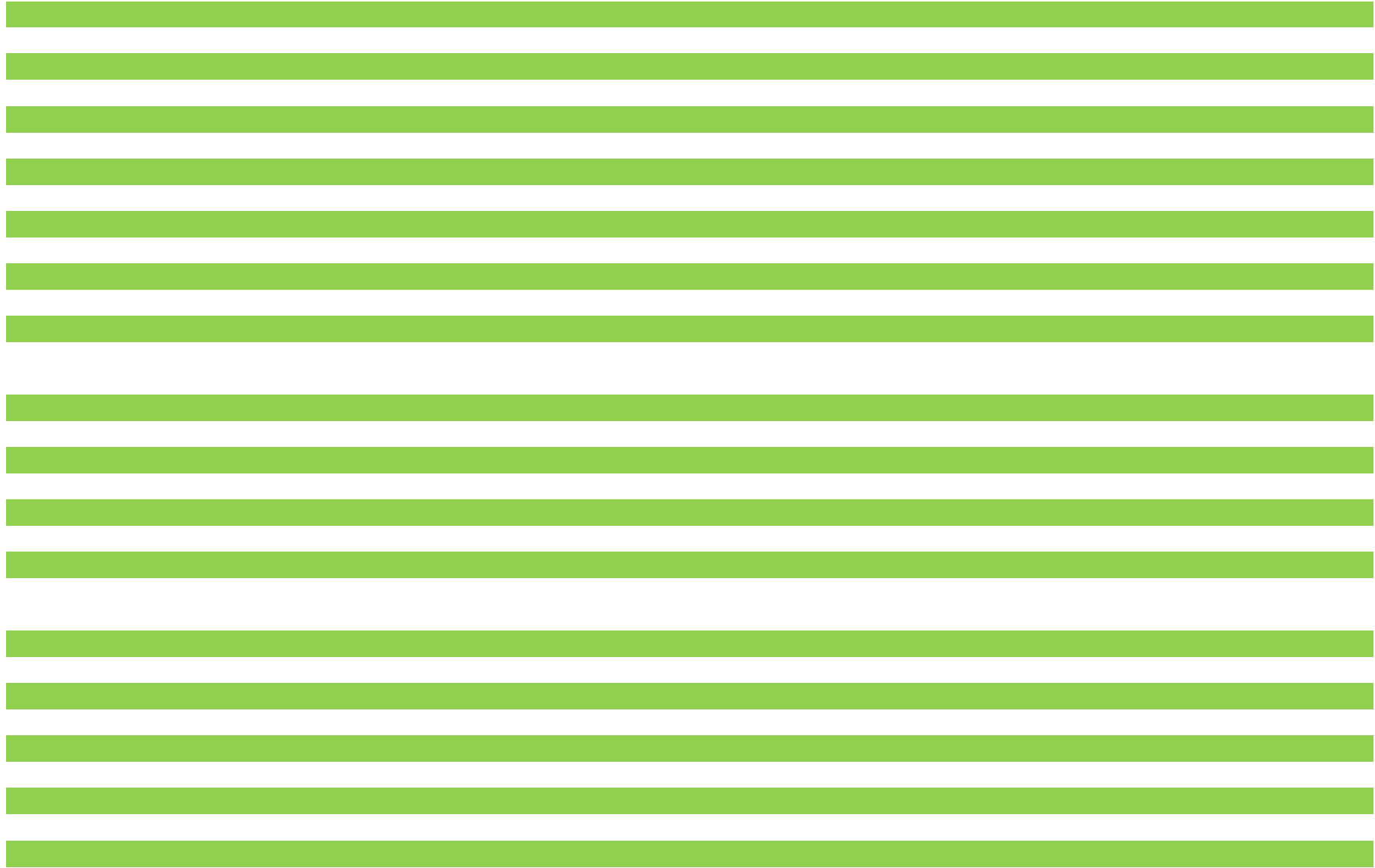
TAB]



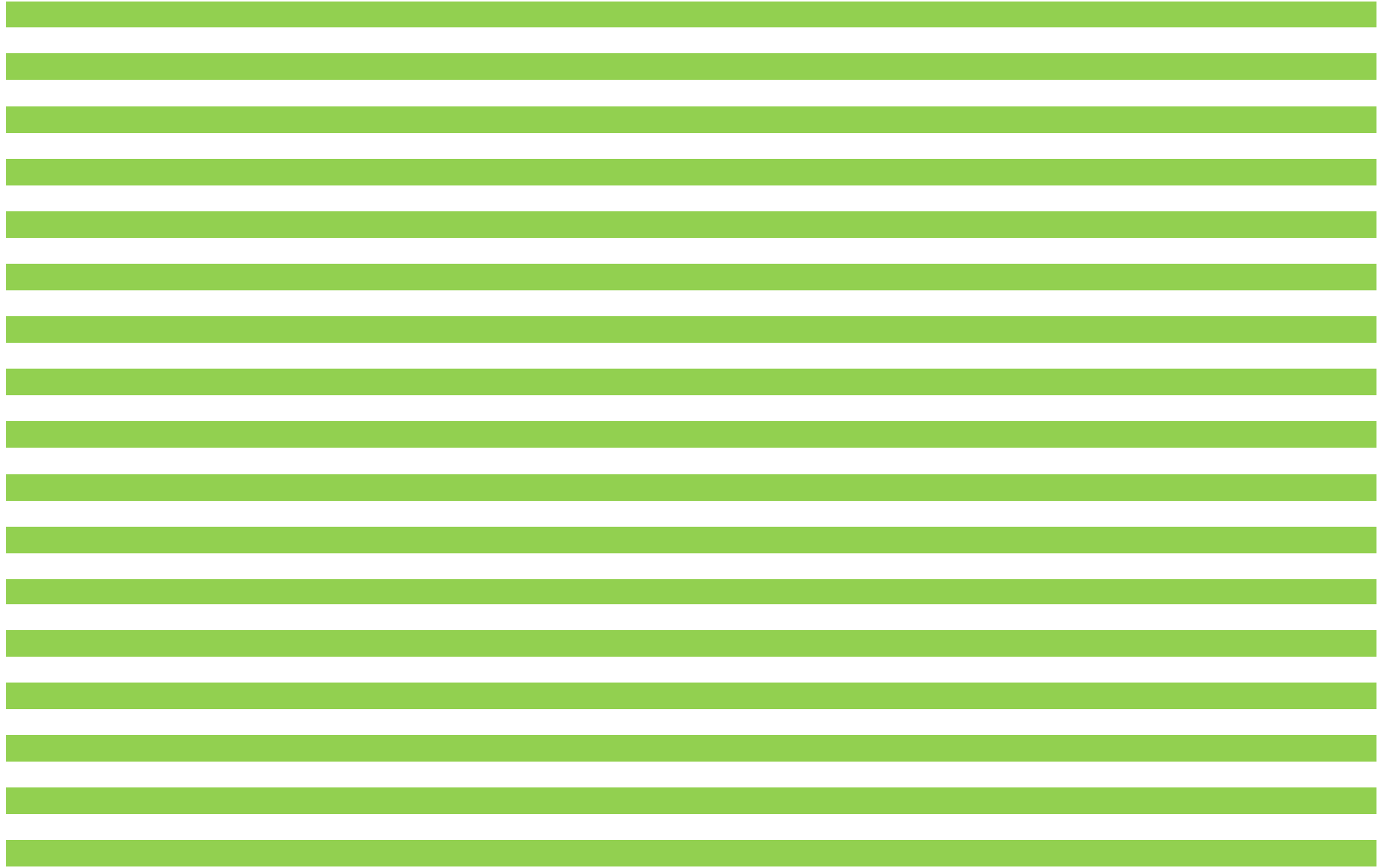
TAB]



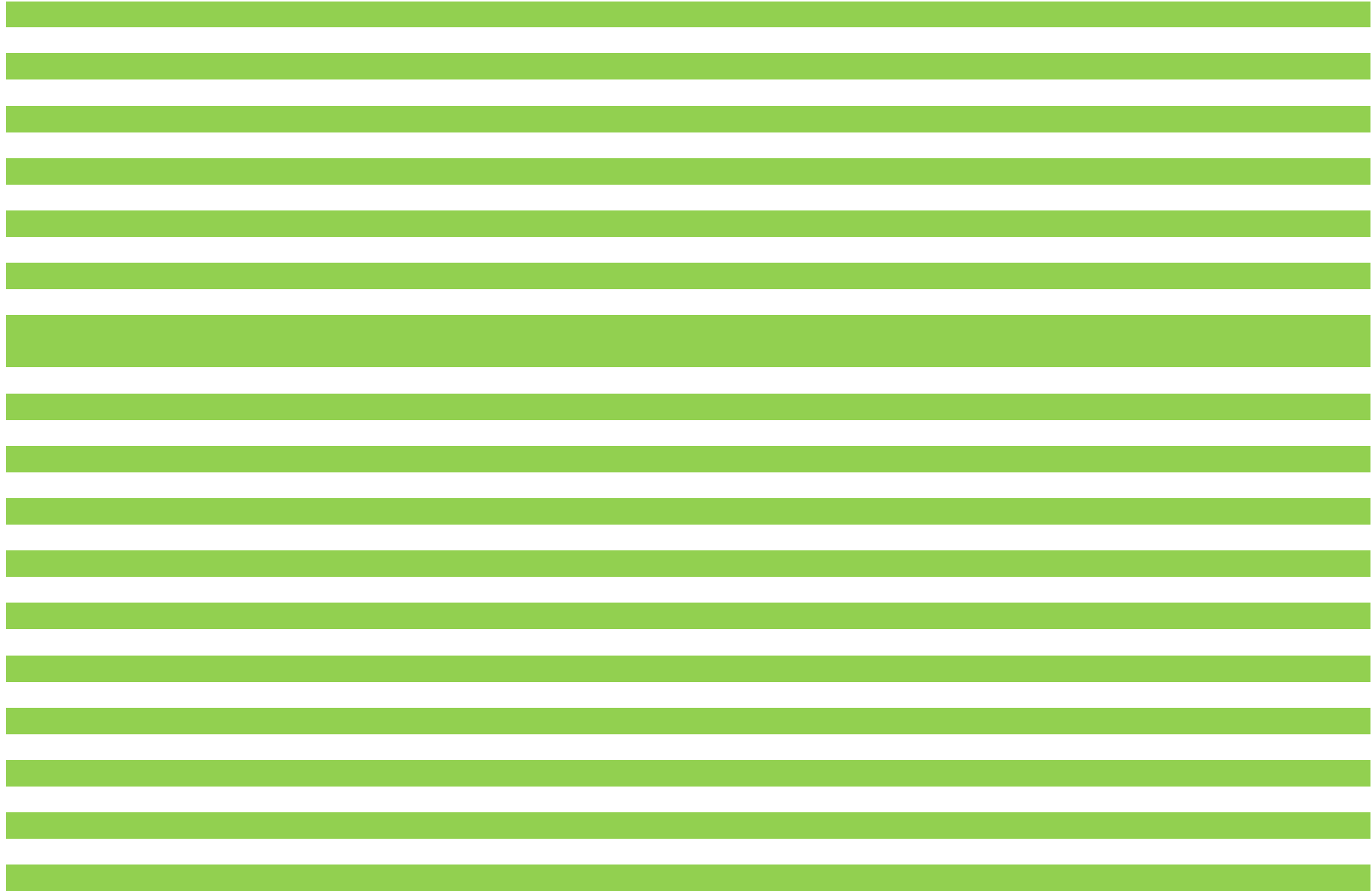
TAB]



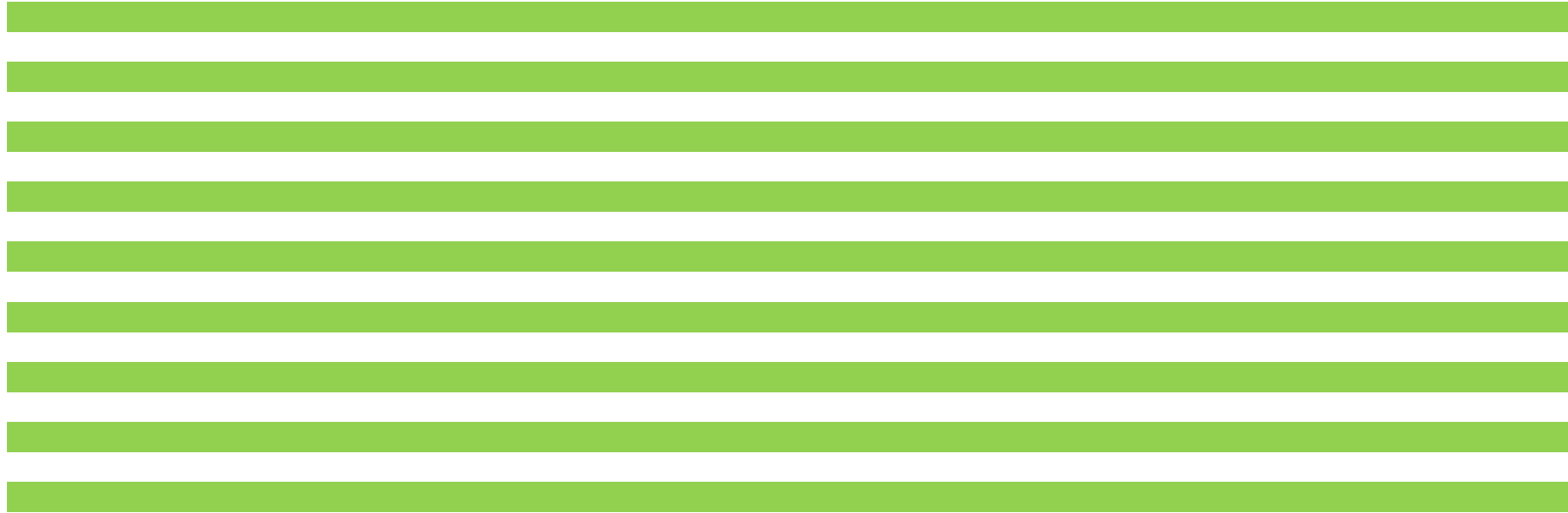
TAB]



TAB]



TAB]



TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

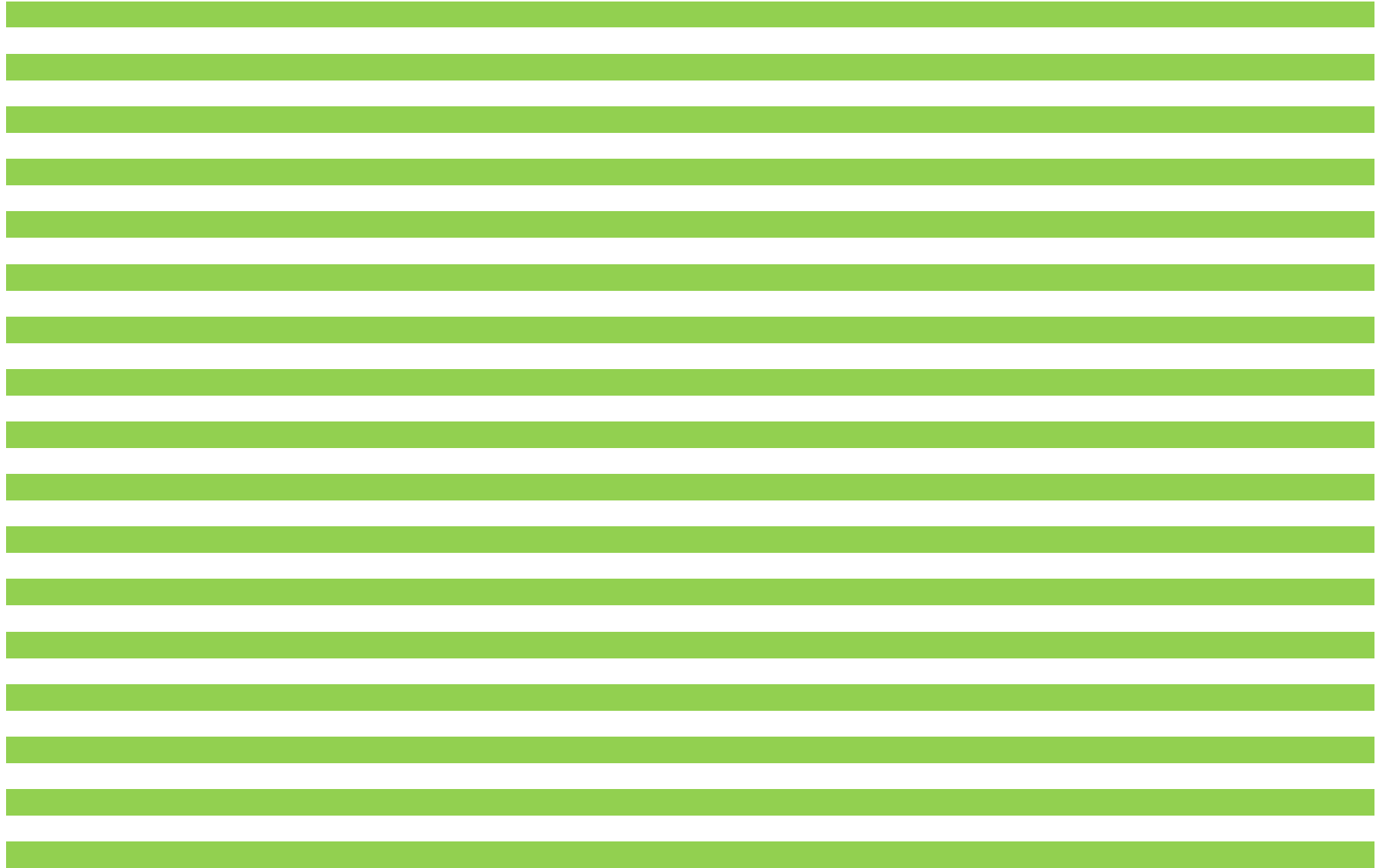
TAB]

TAB]

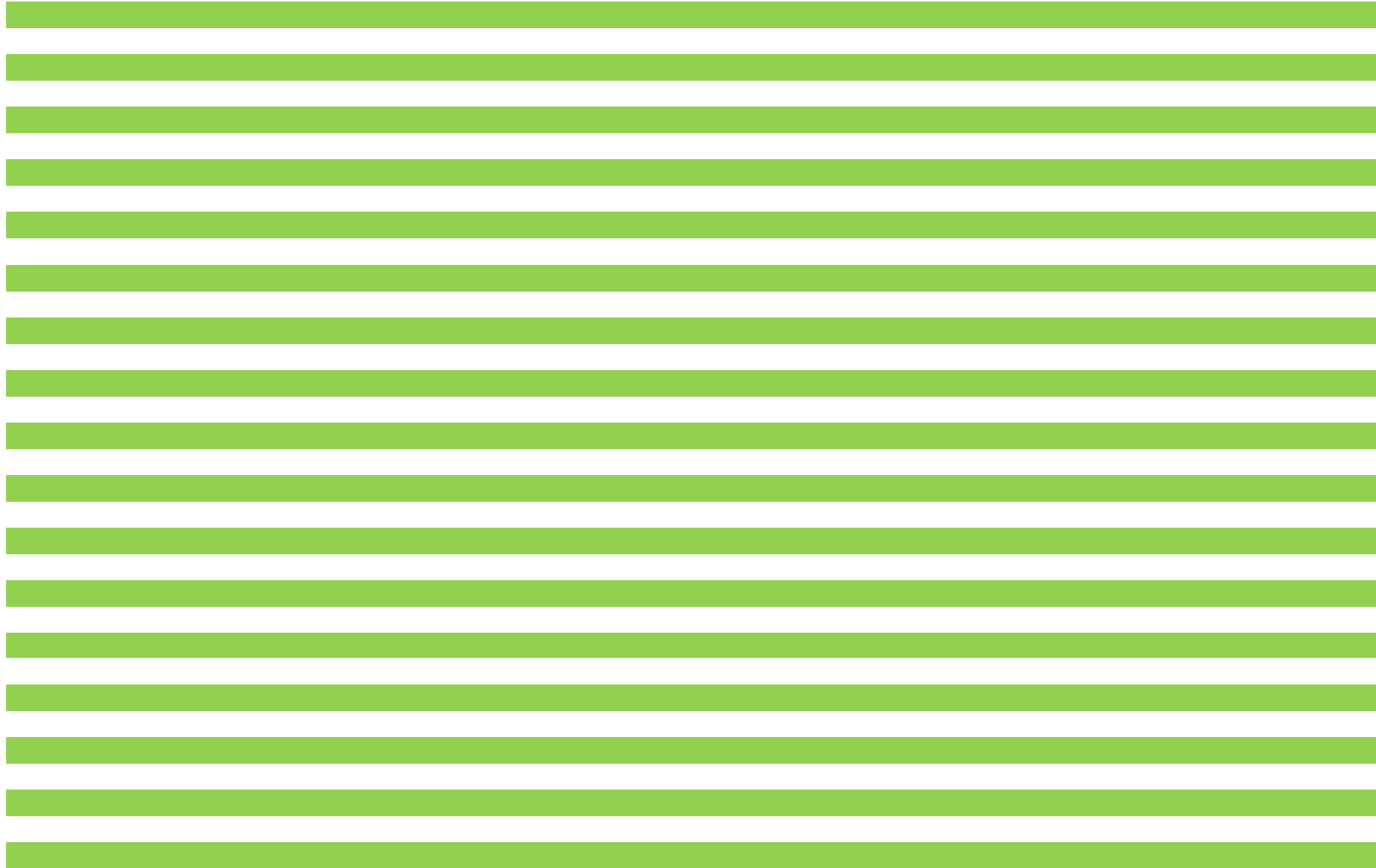
TAB]

TAB]

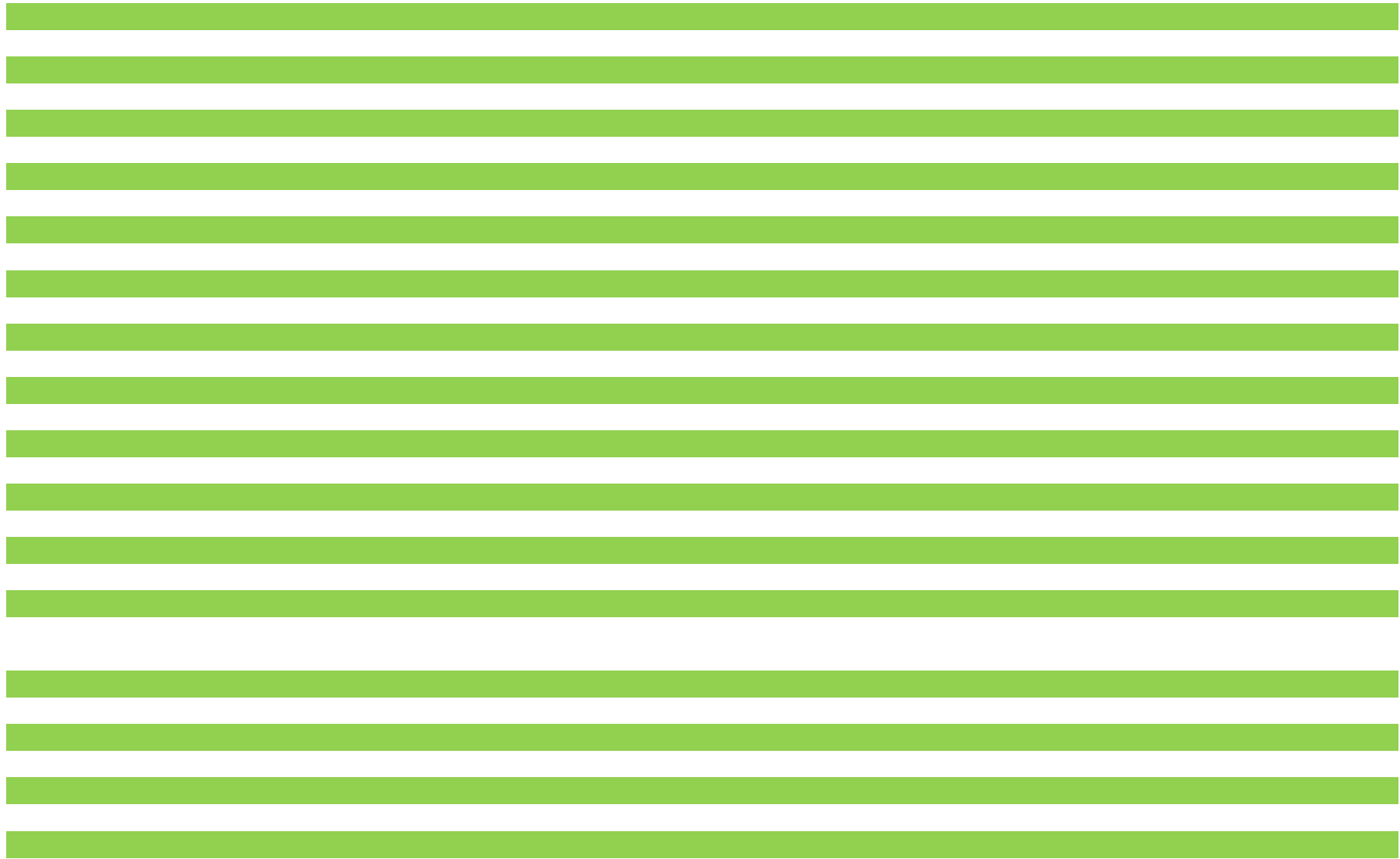
TAB]



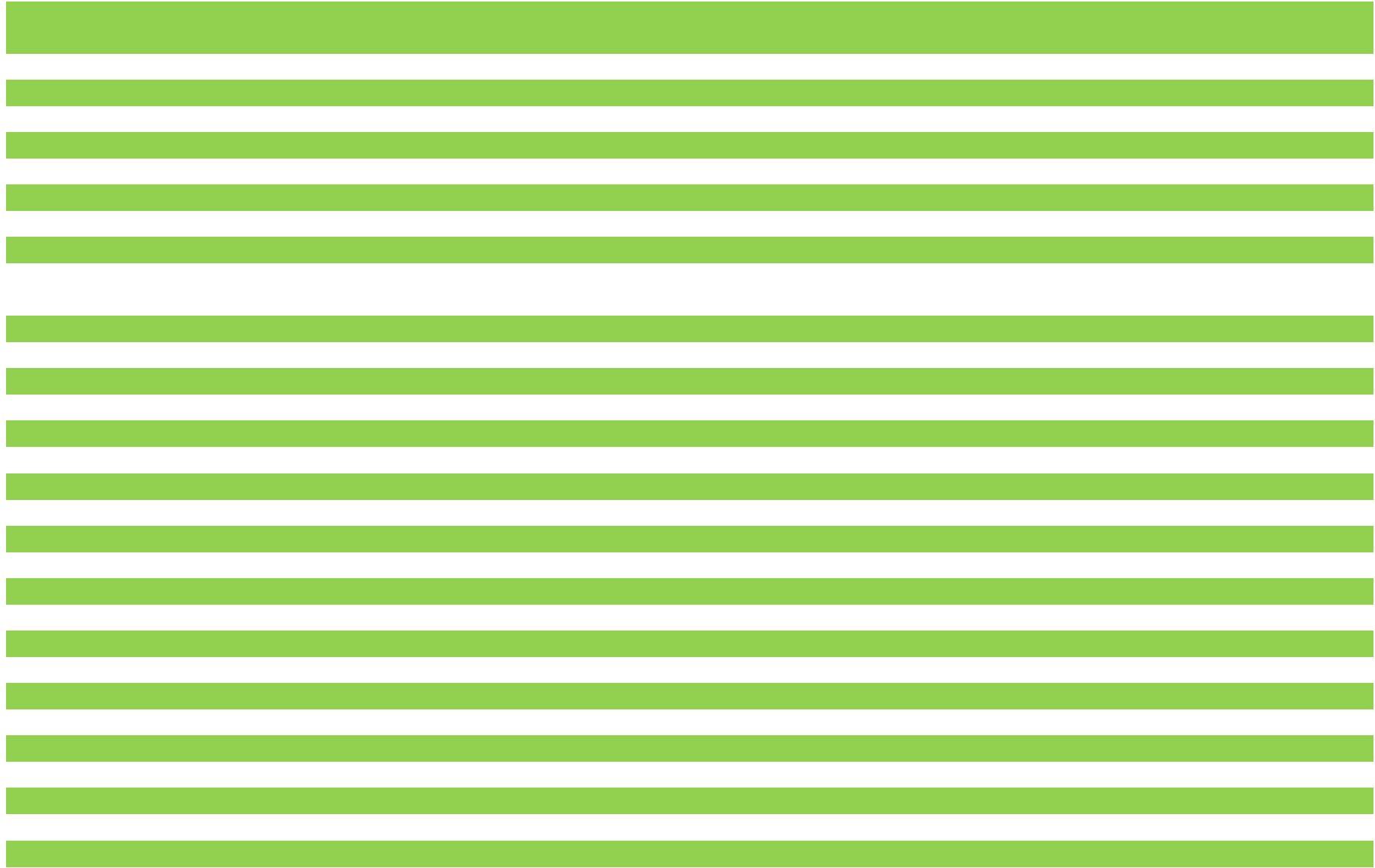
TAB]



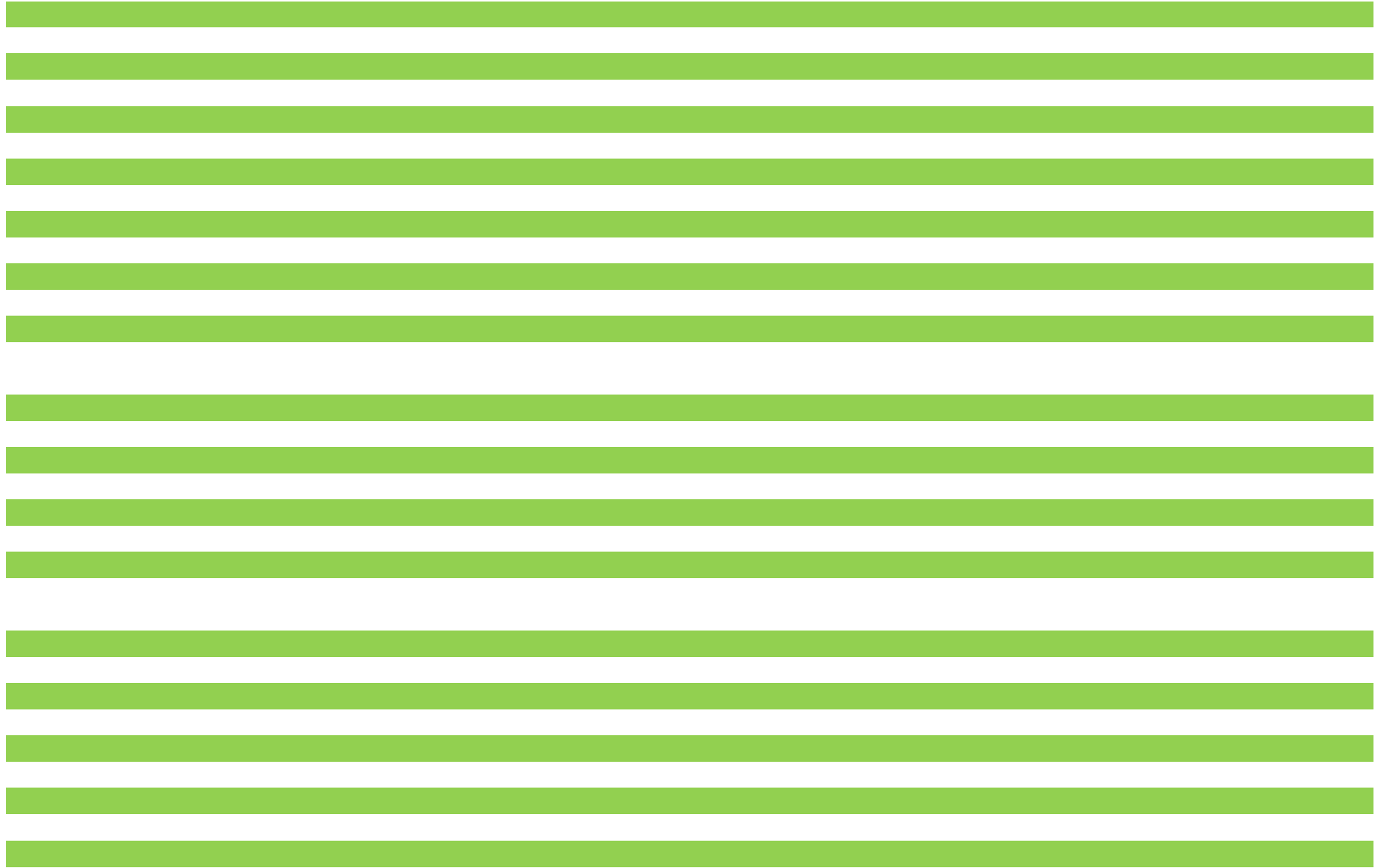
TAB]



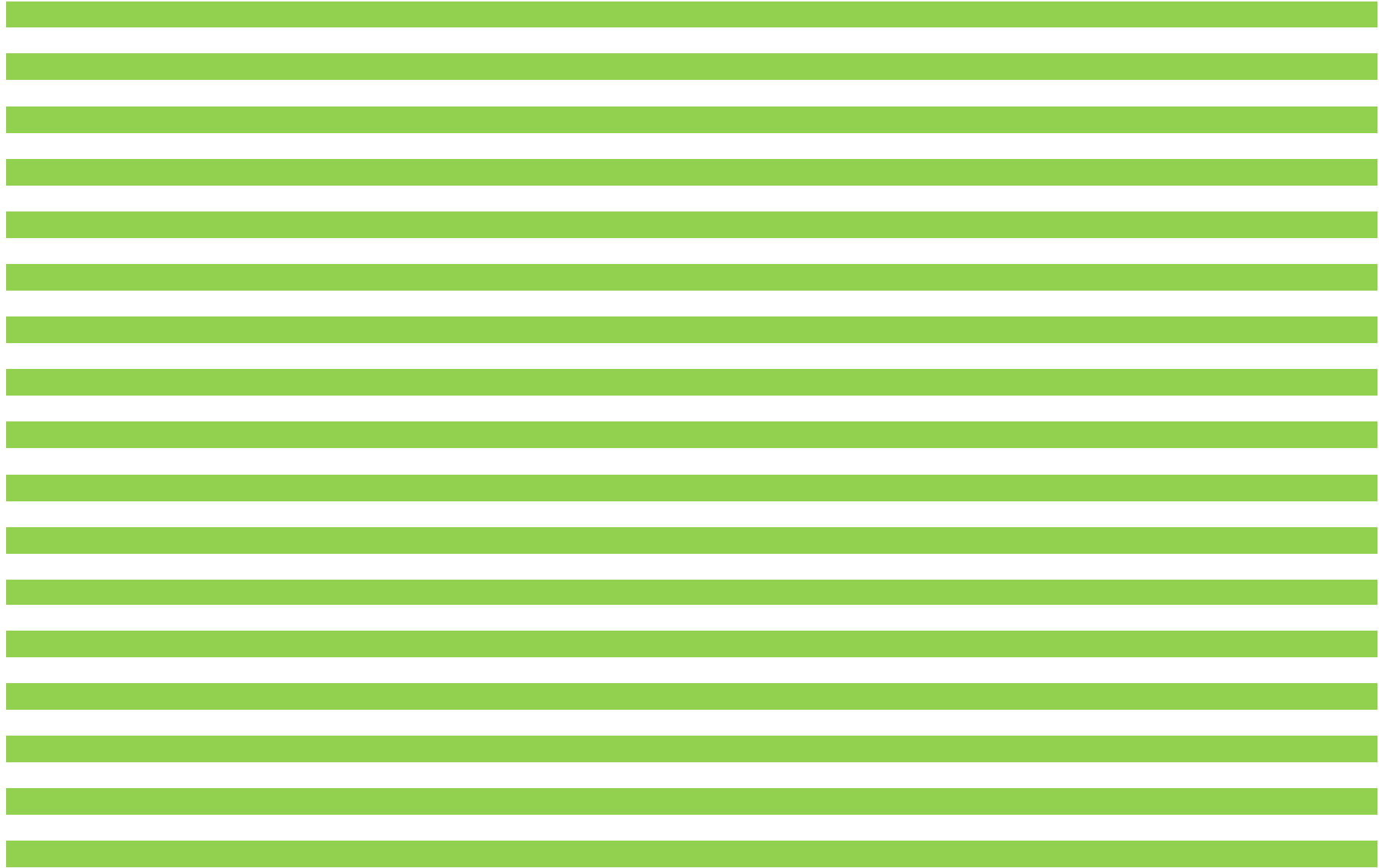
TAB]



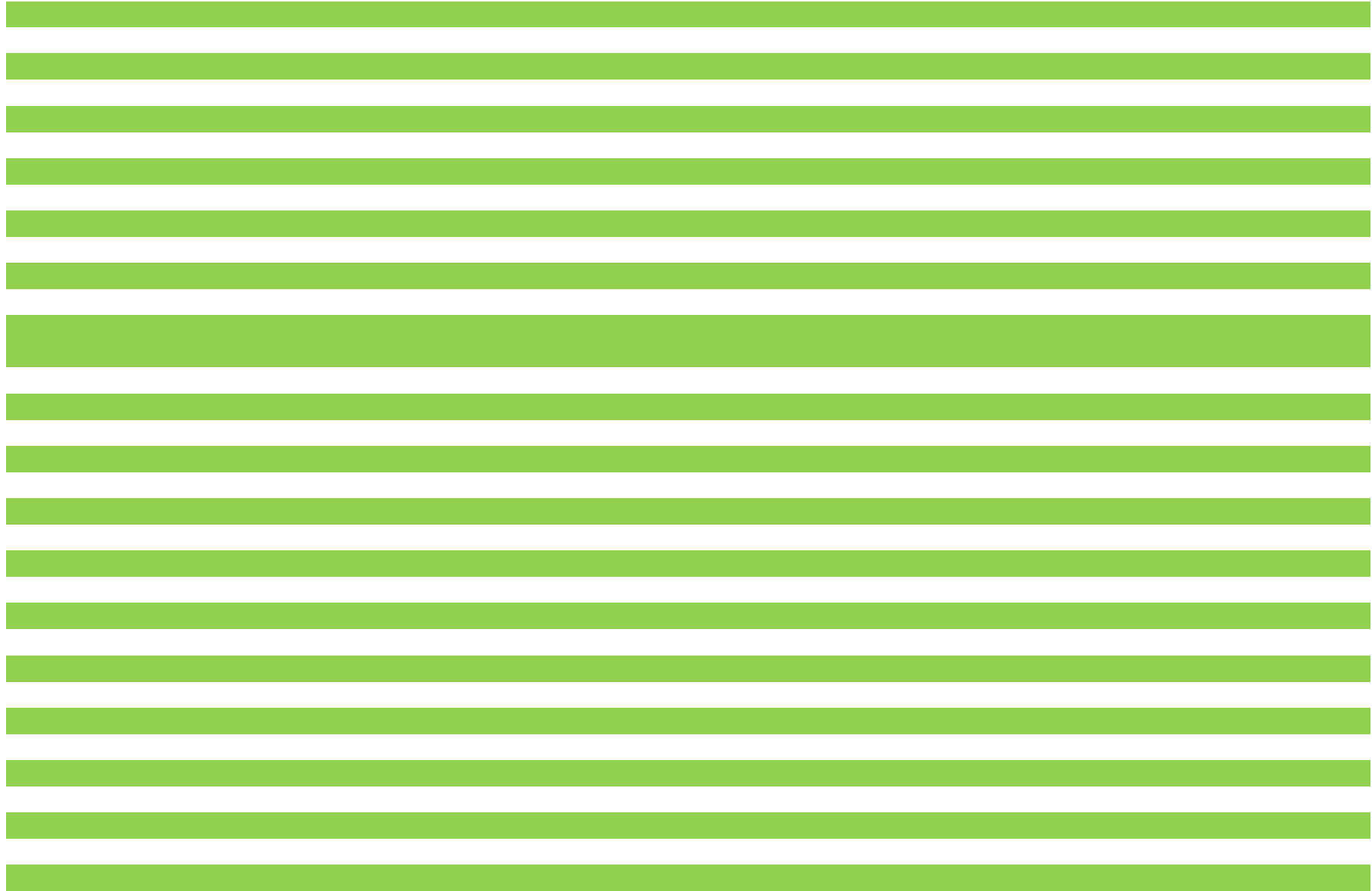
TAB]



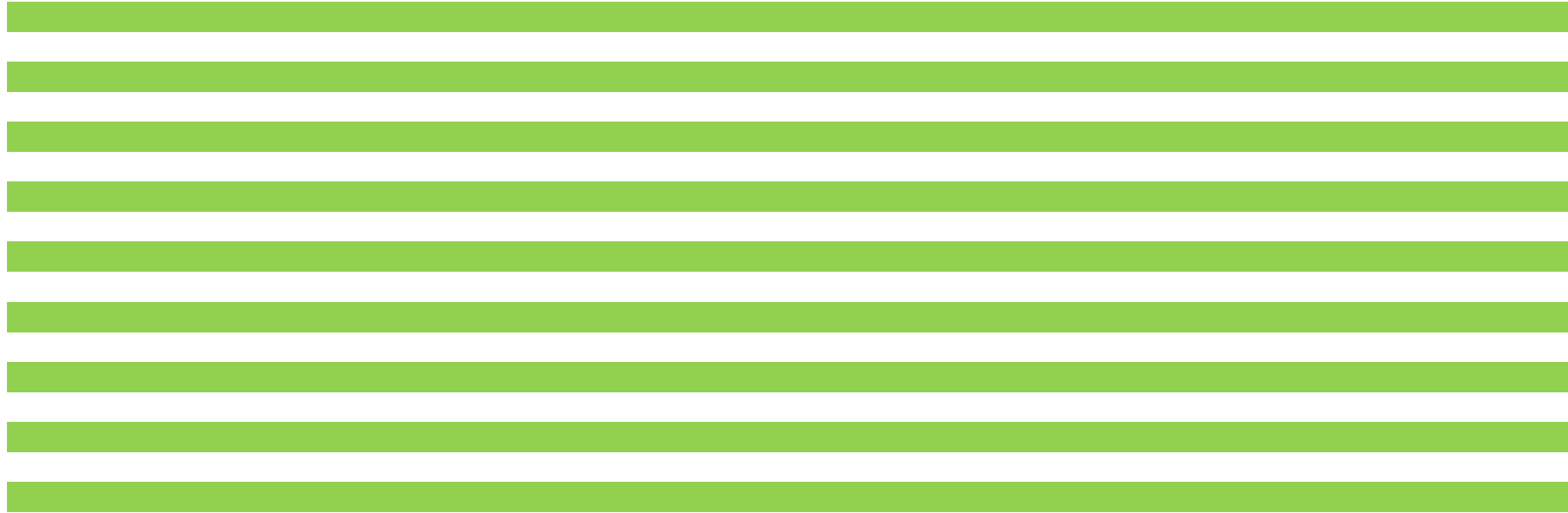
TAB]



TAB]



TAB]



TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

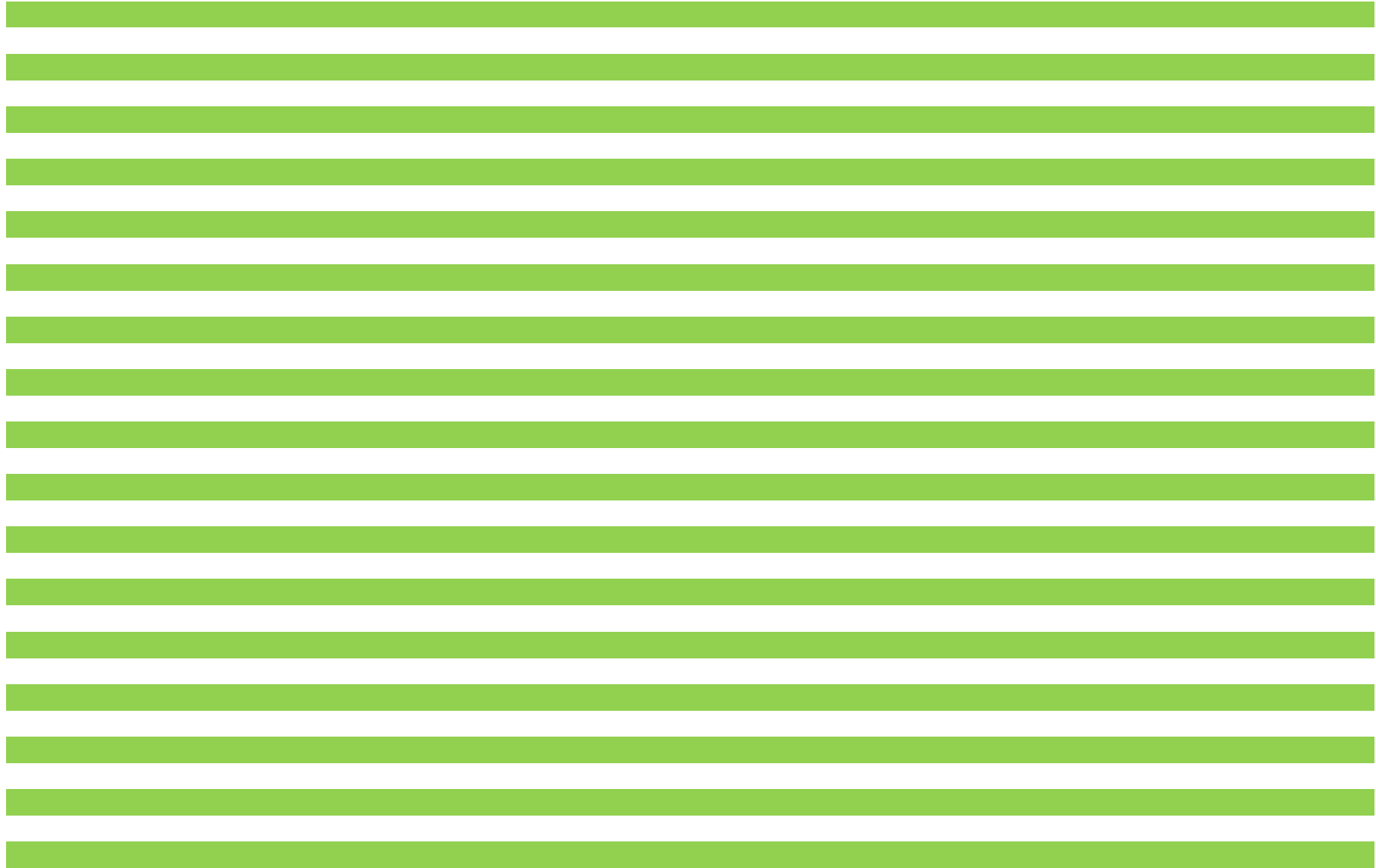
TAB]

TAB]

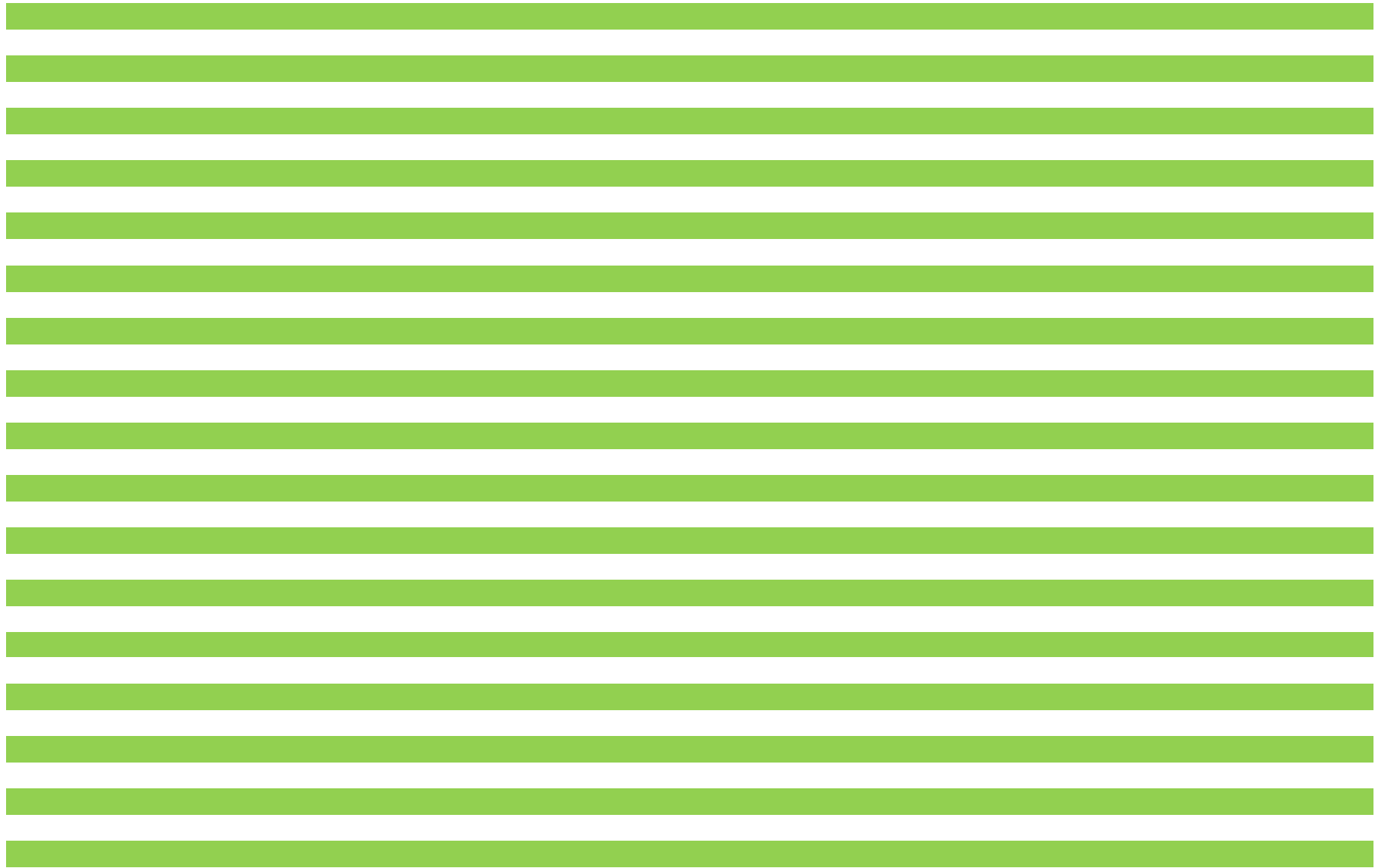
TAB]

TAB]

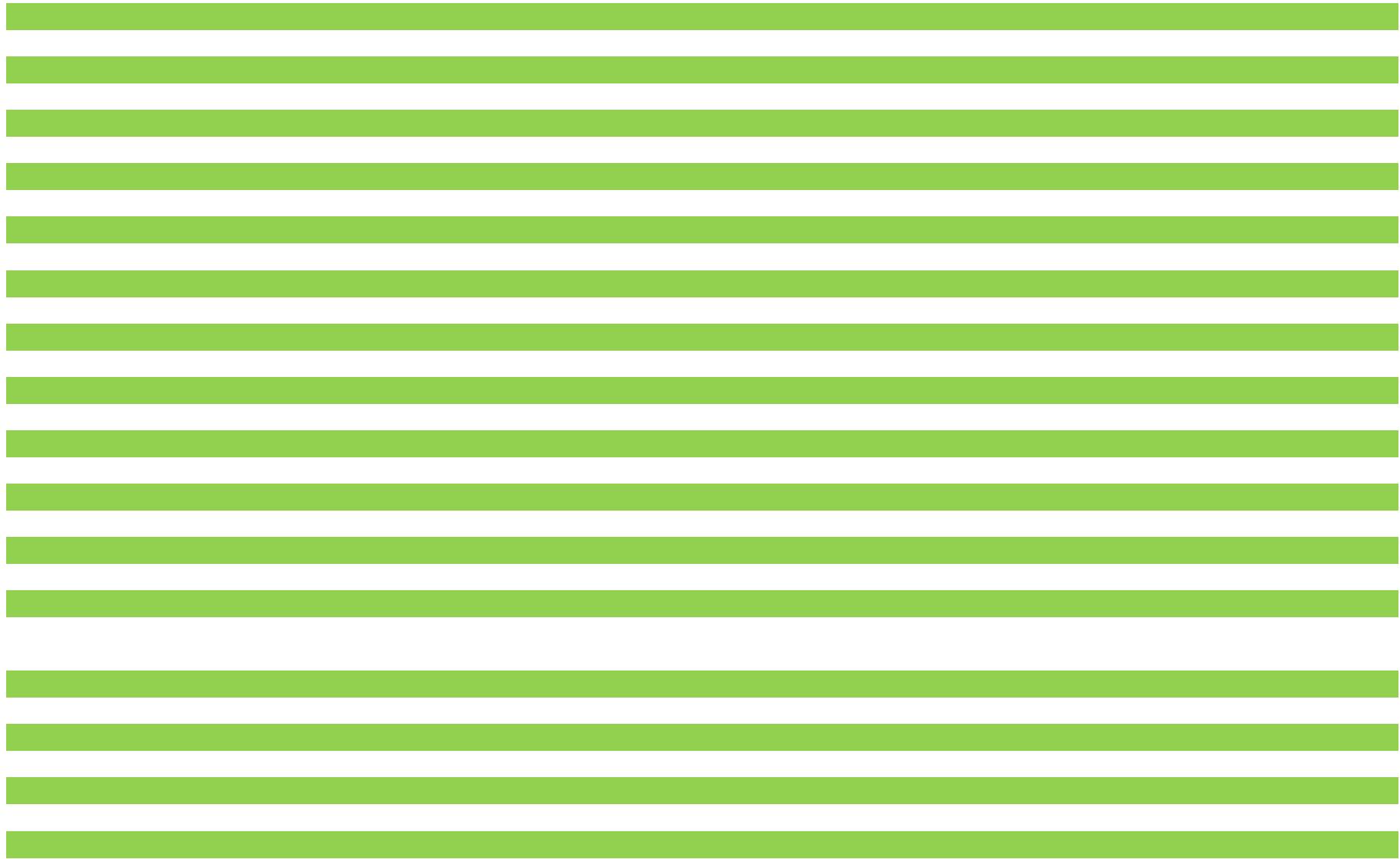
TAB]



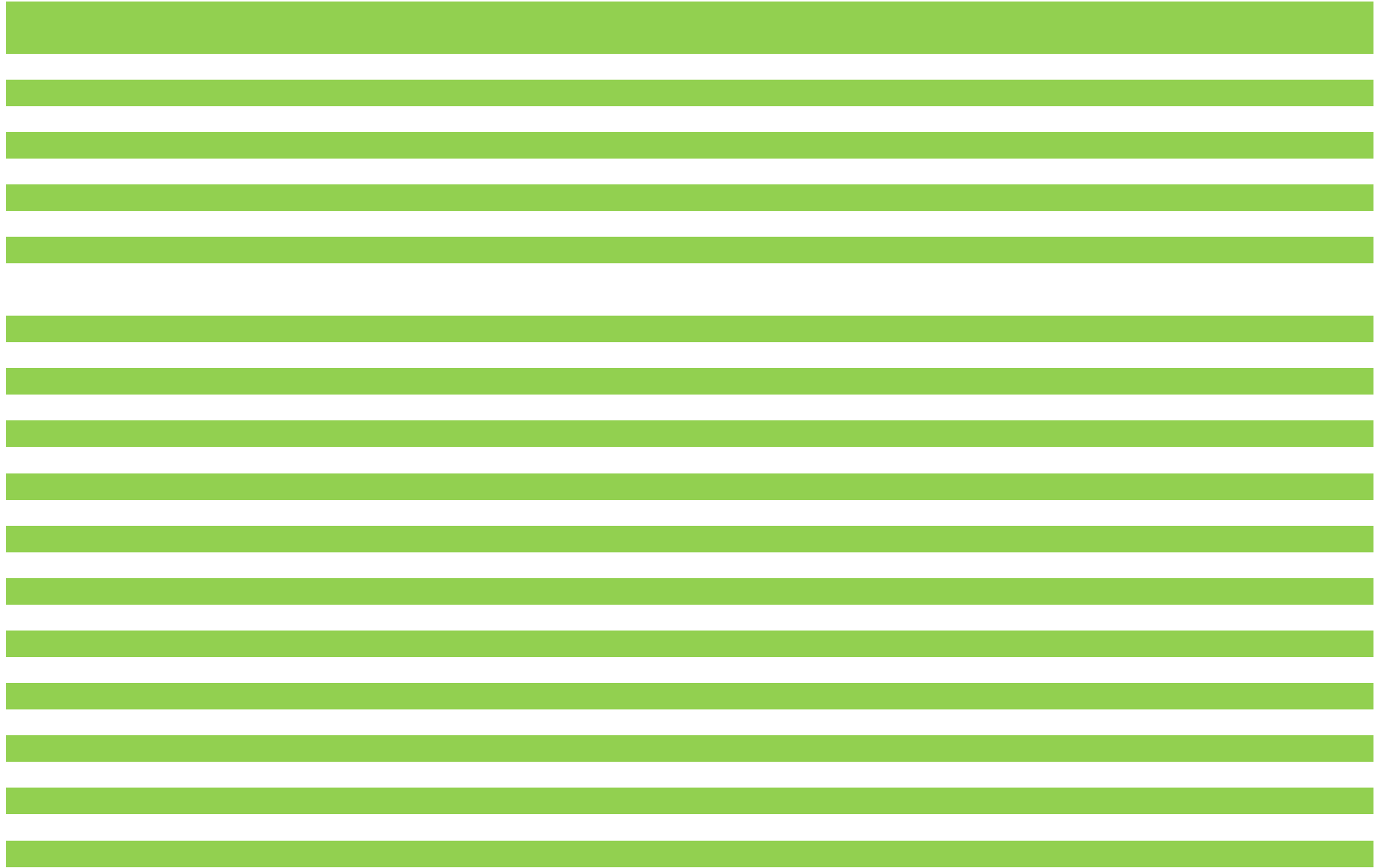
TAB]



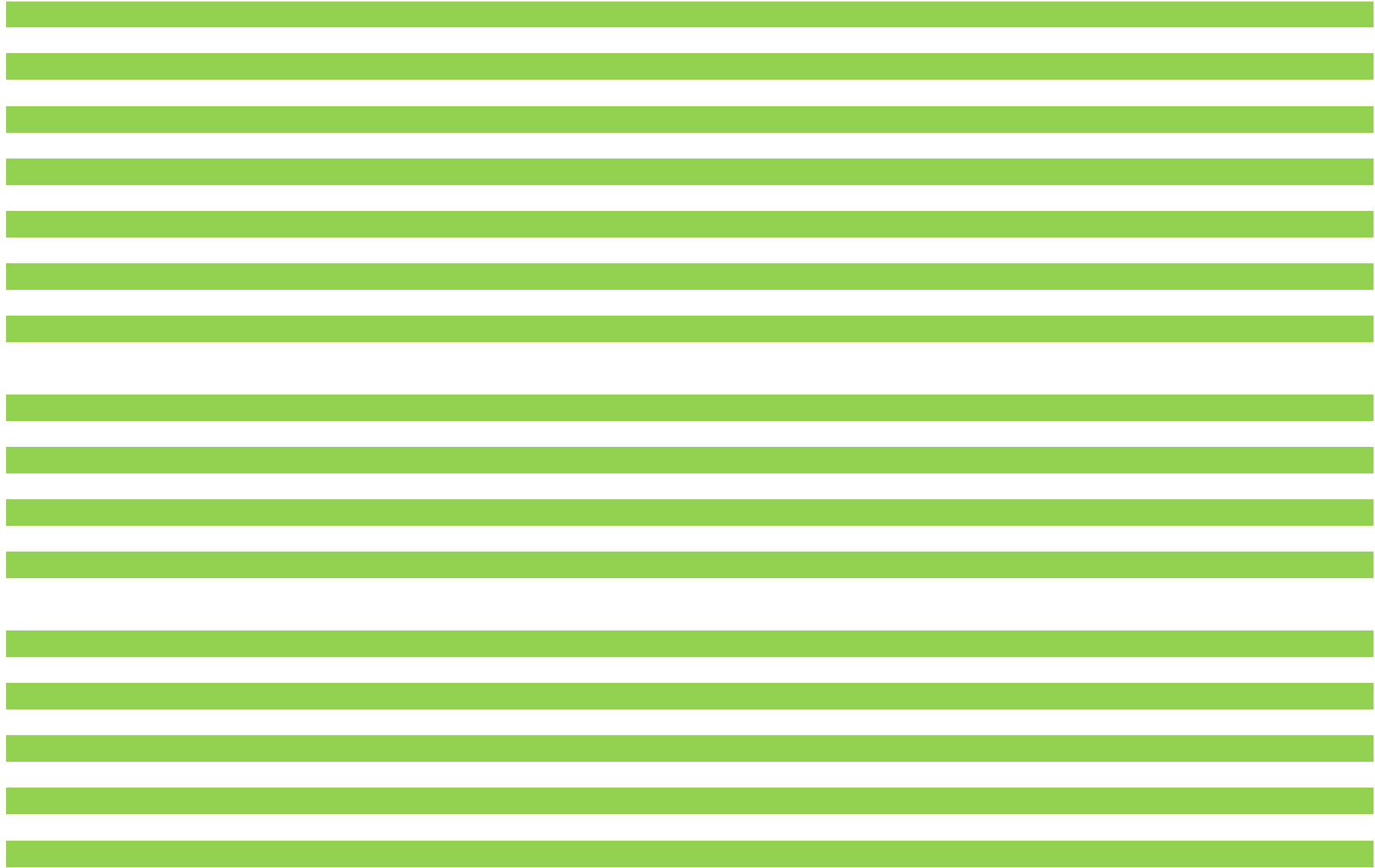
TAB]



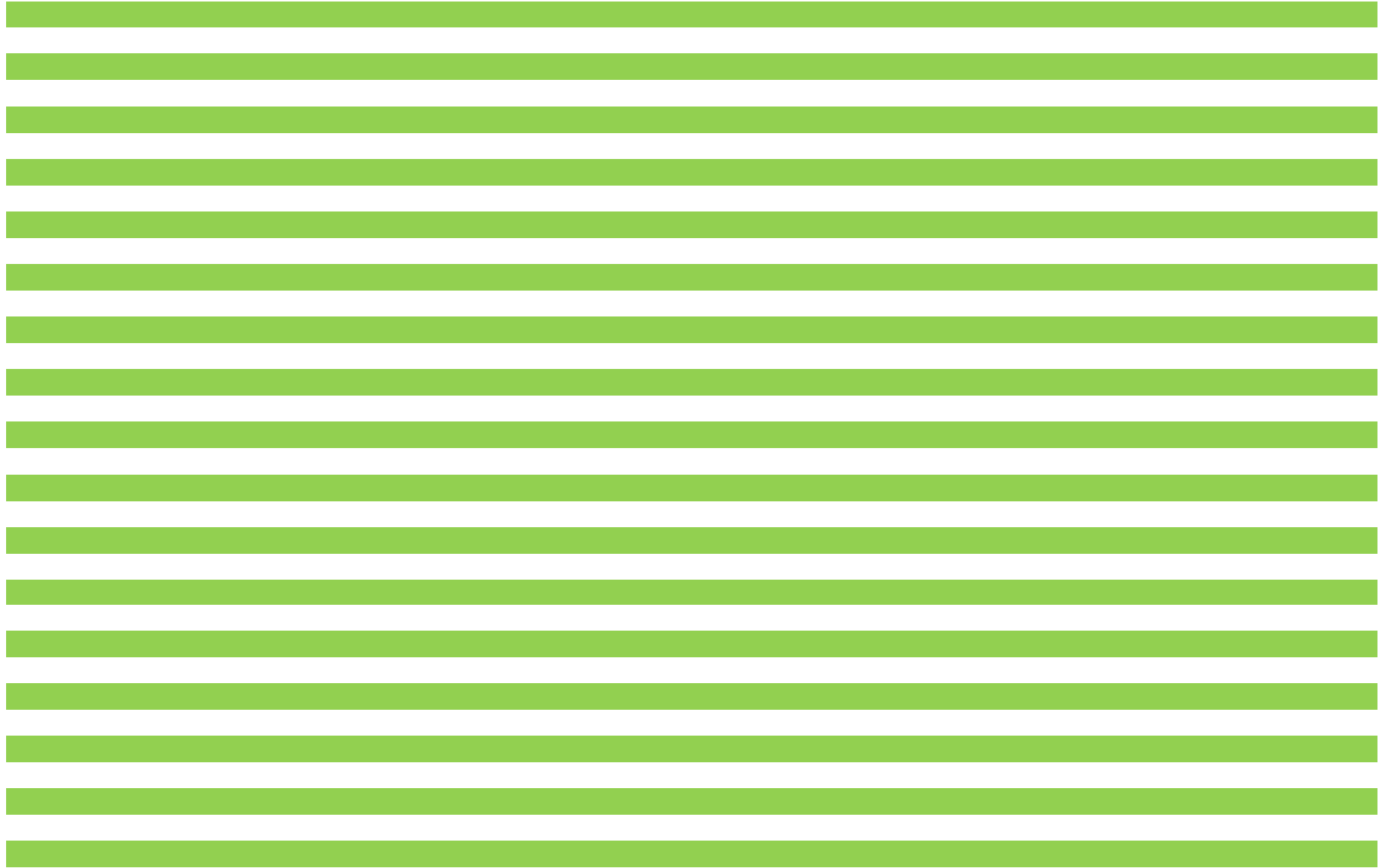
TAB]



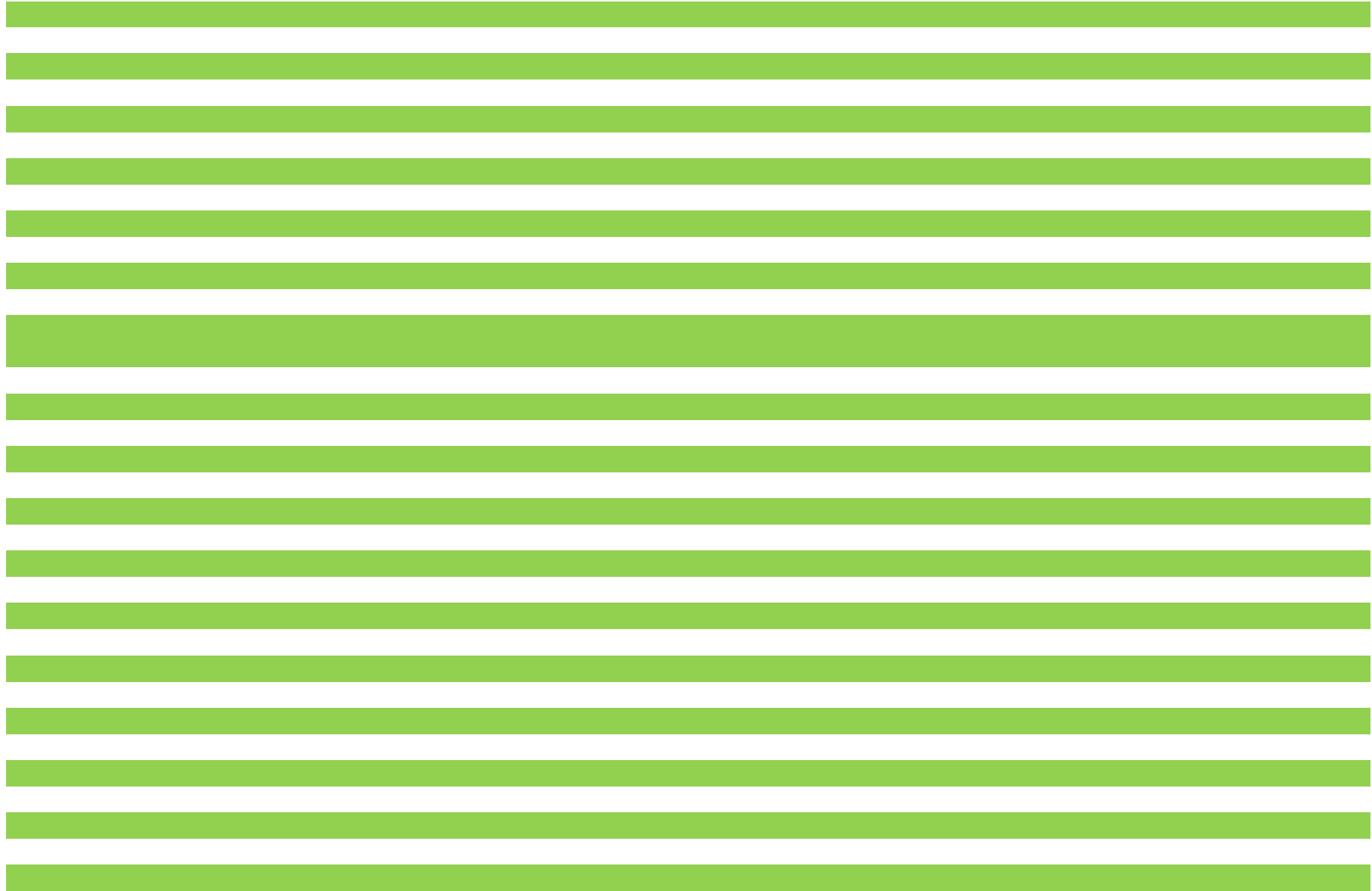
TAB]



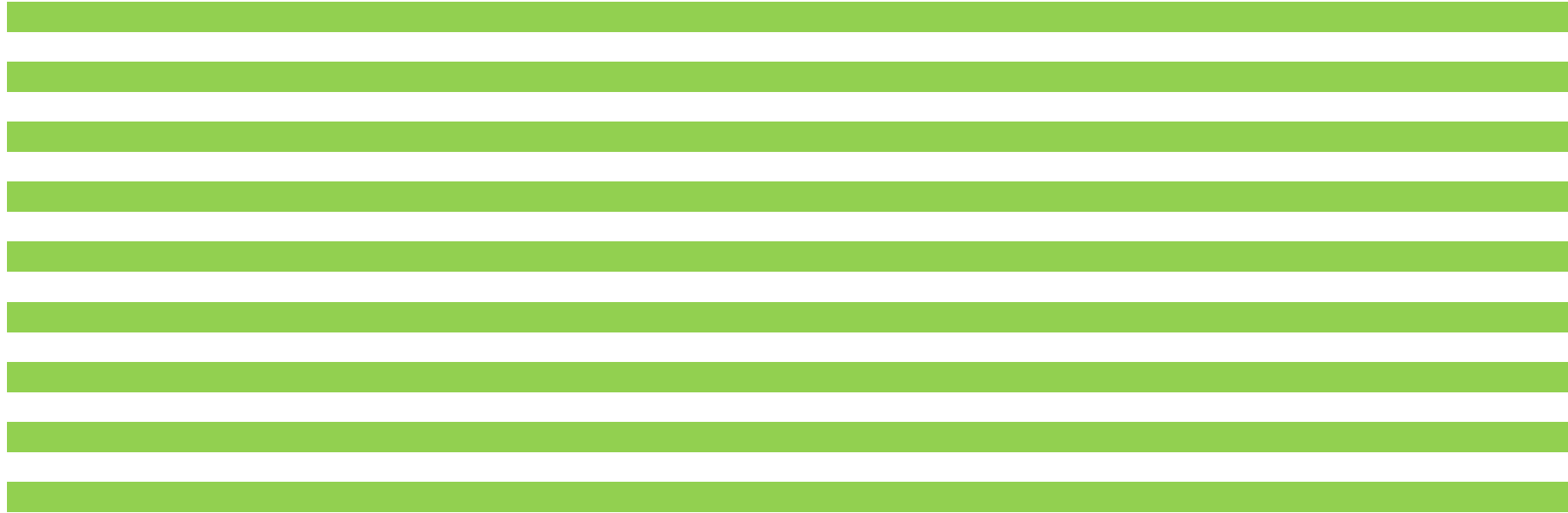
TAB]



TAB]



TAB]



TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

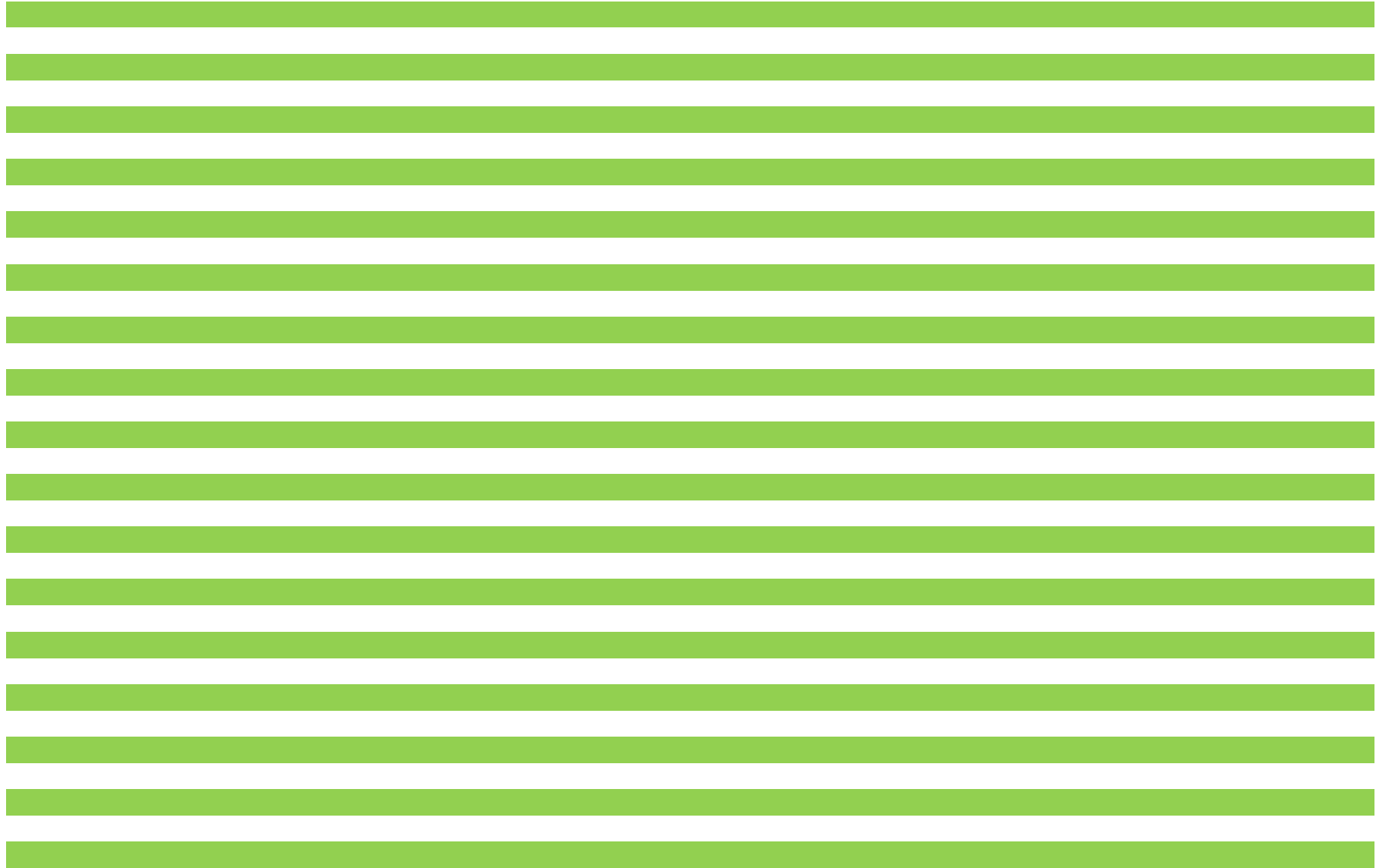
TAB]

TAB]

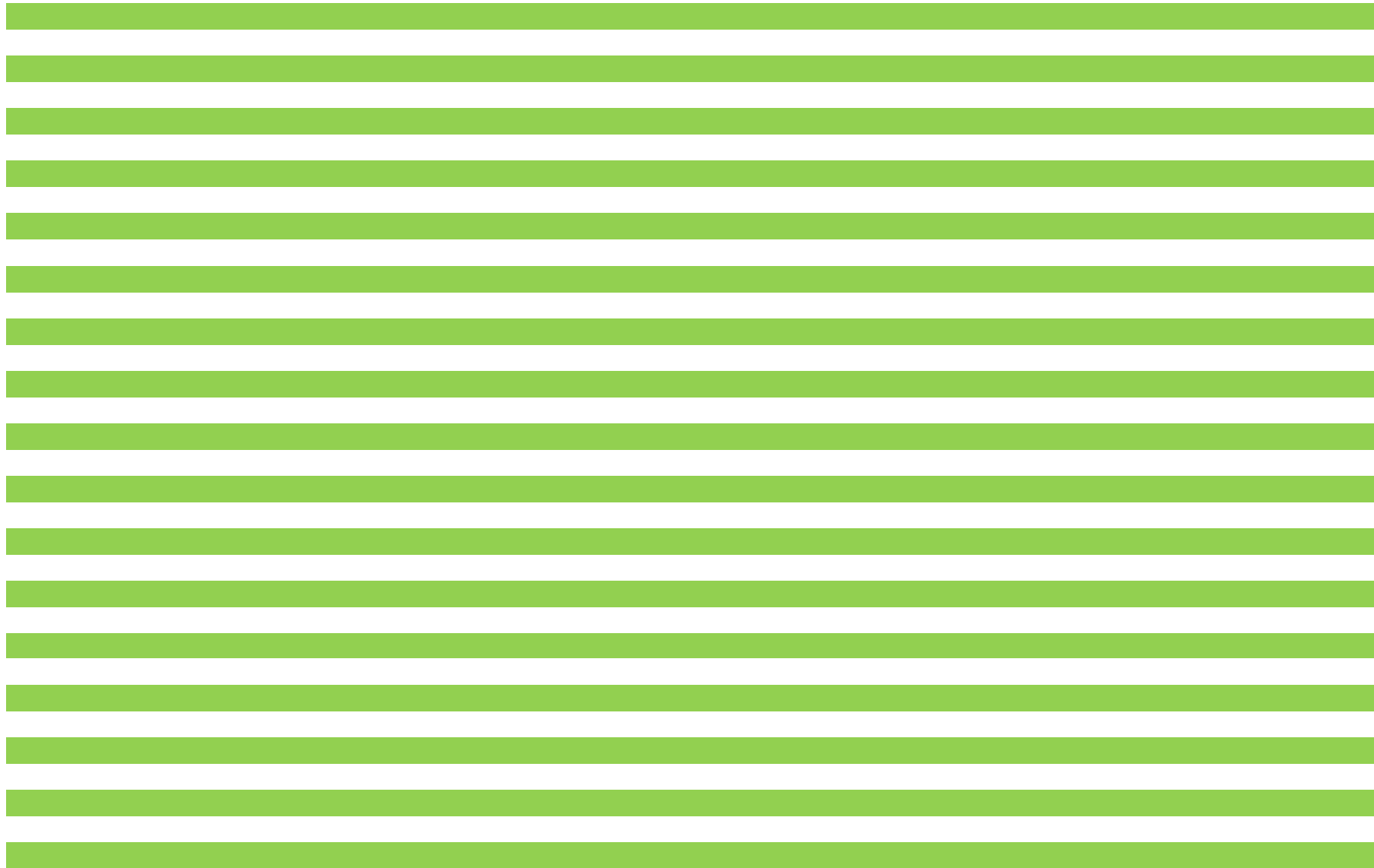
TAB]

TAB]

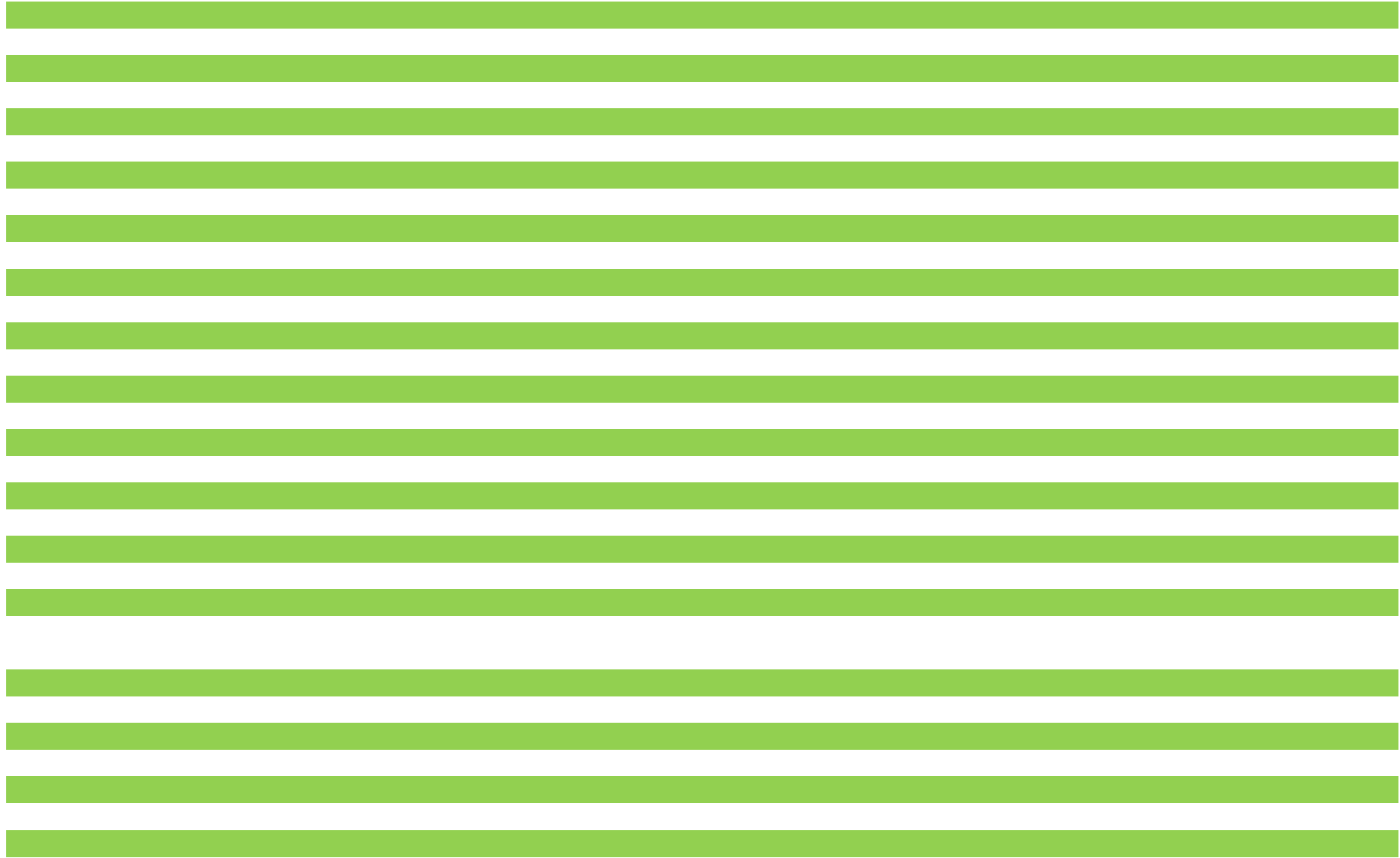
TAB]



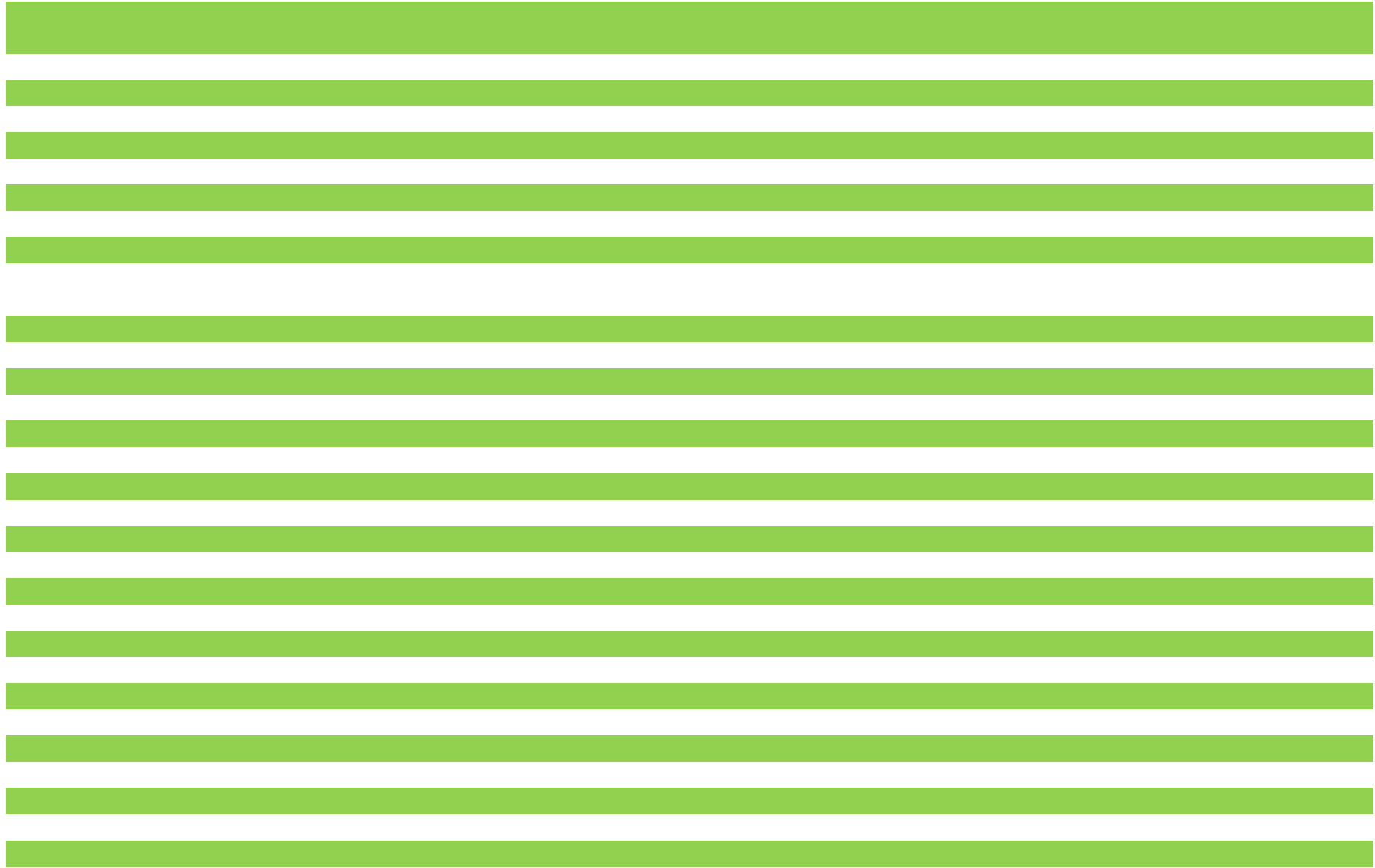
TAB]



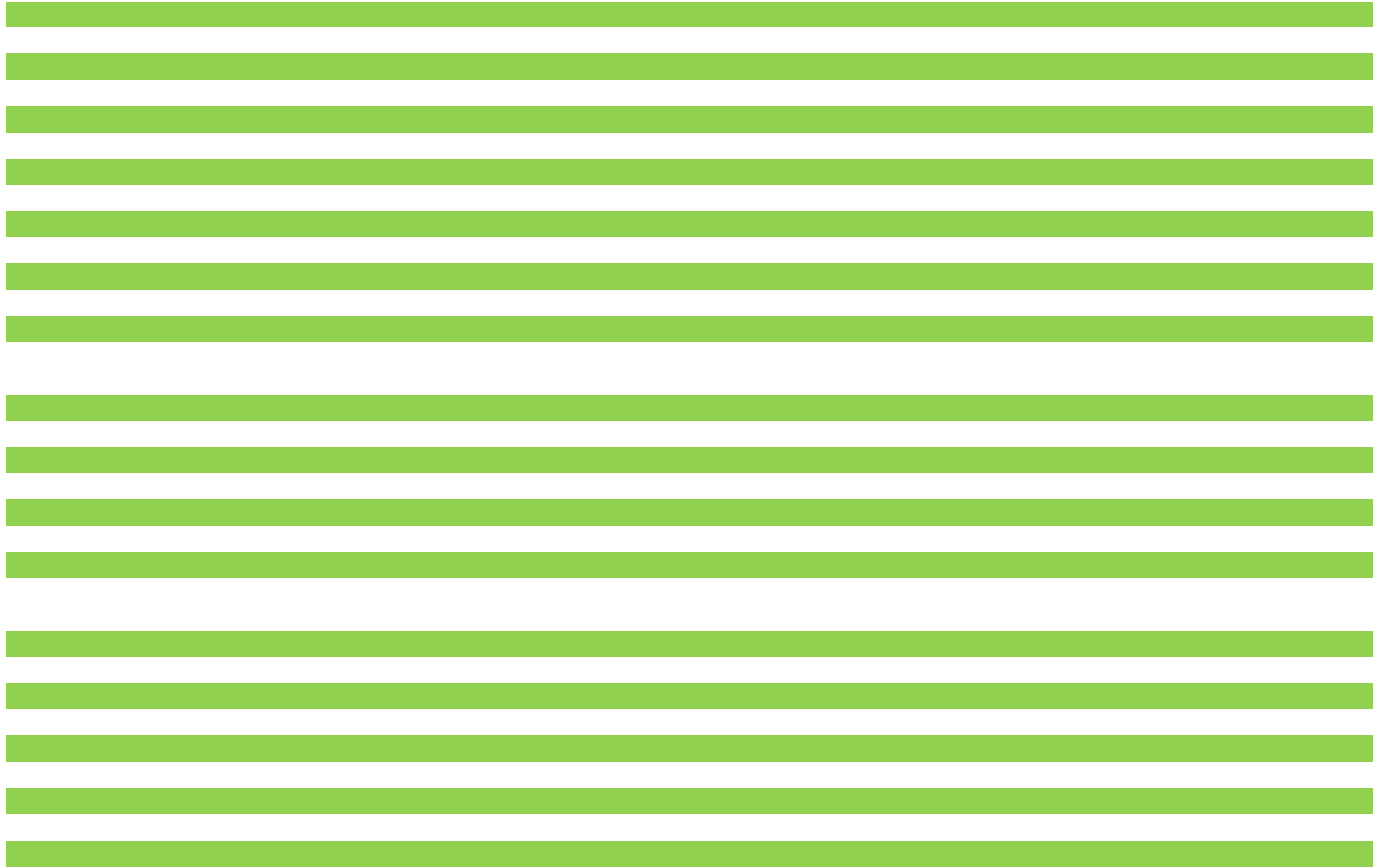
TAB]



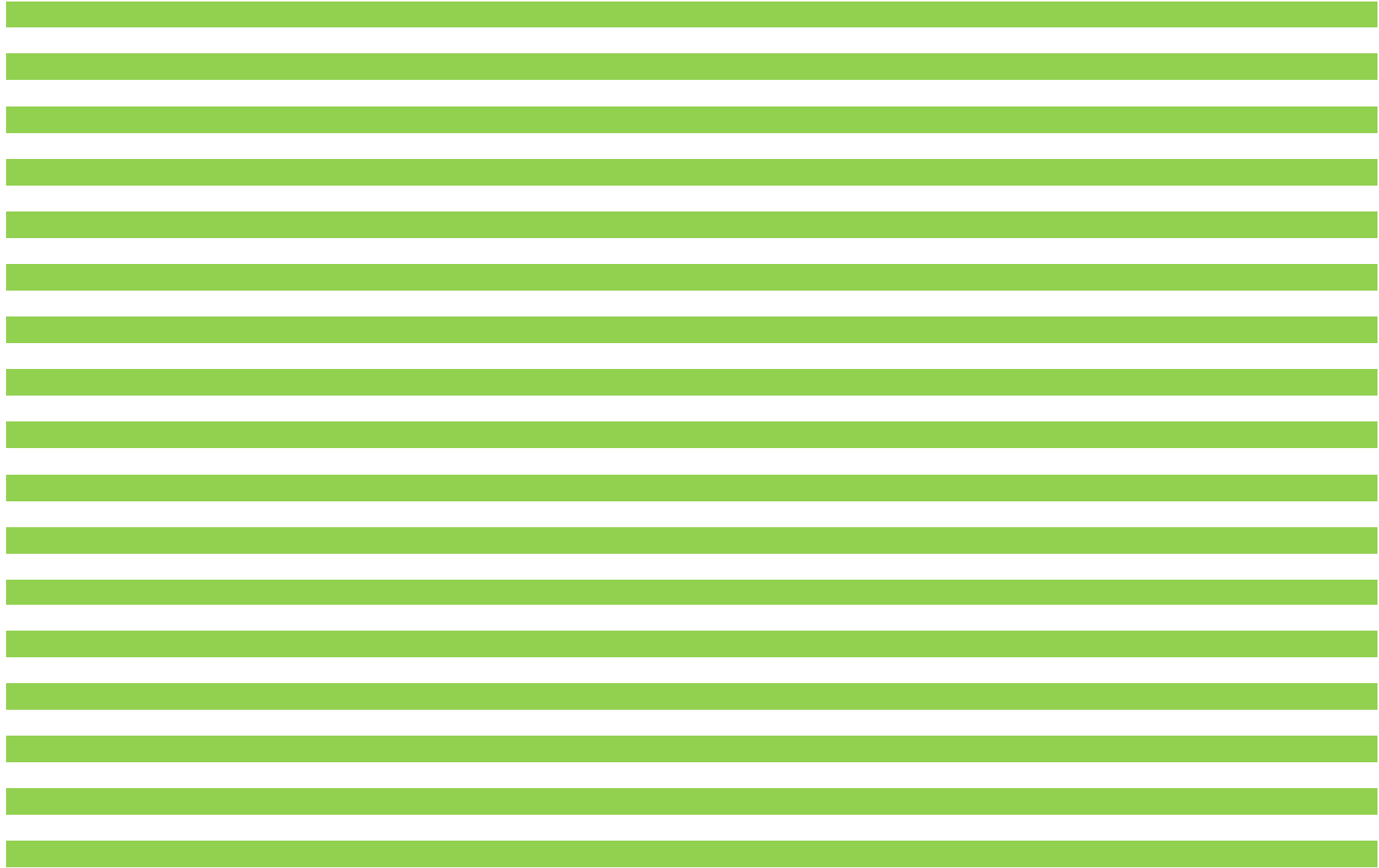
TAB]



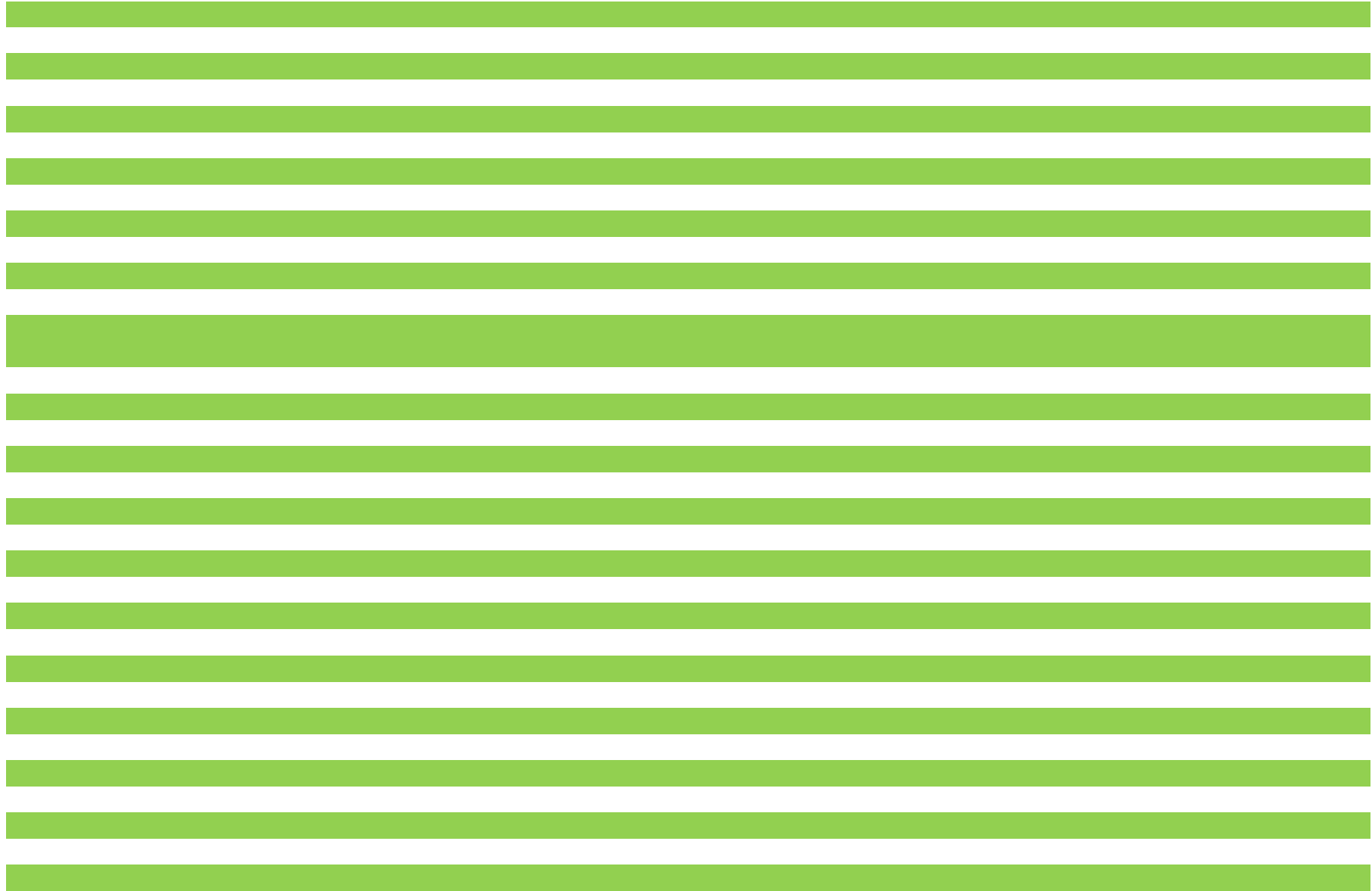
TAB]



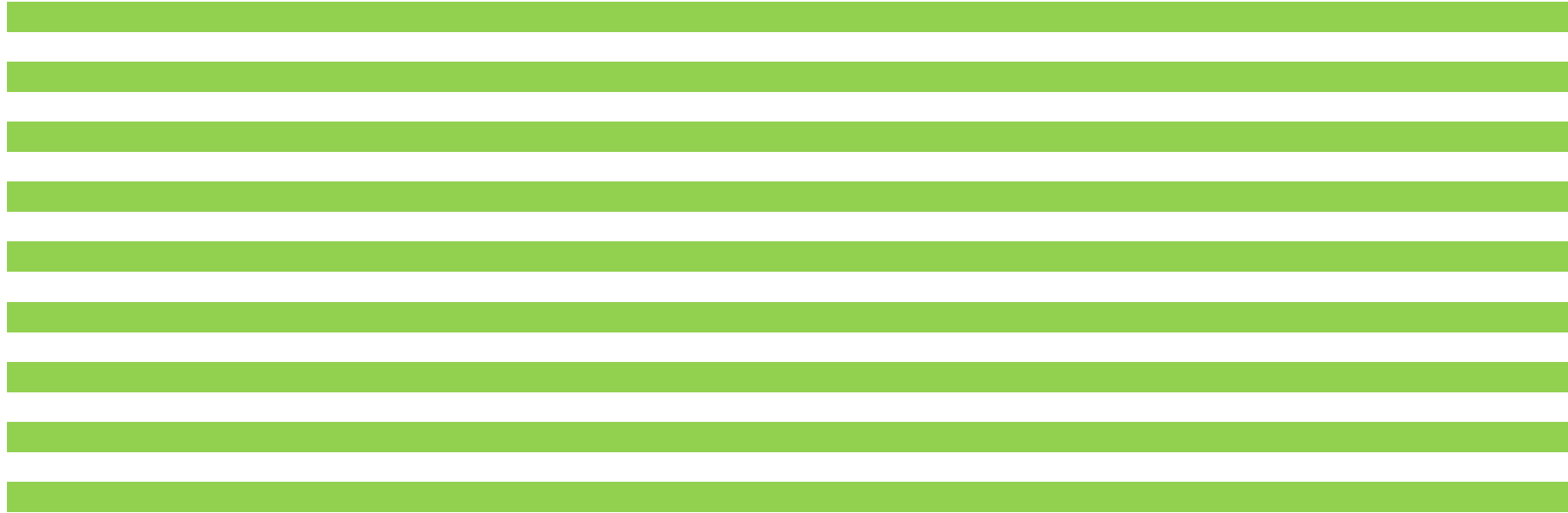
TAB]



TAB]



TAB]



TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

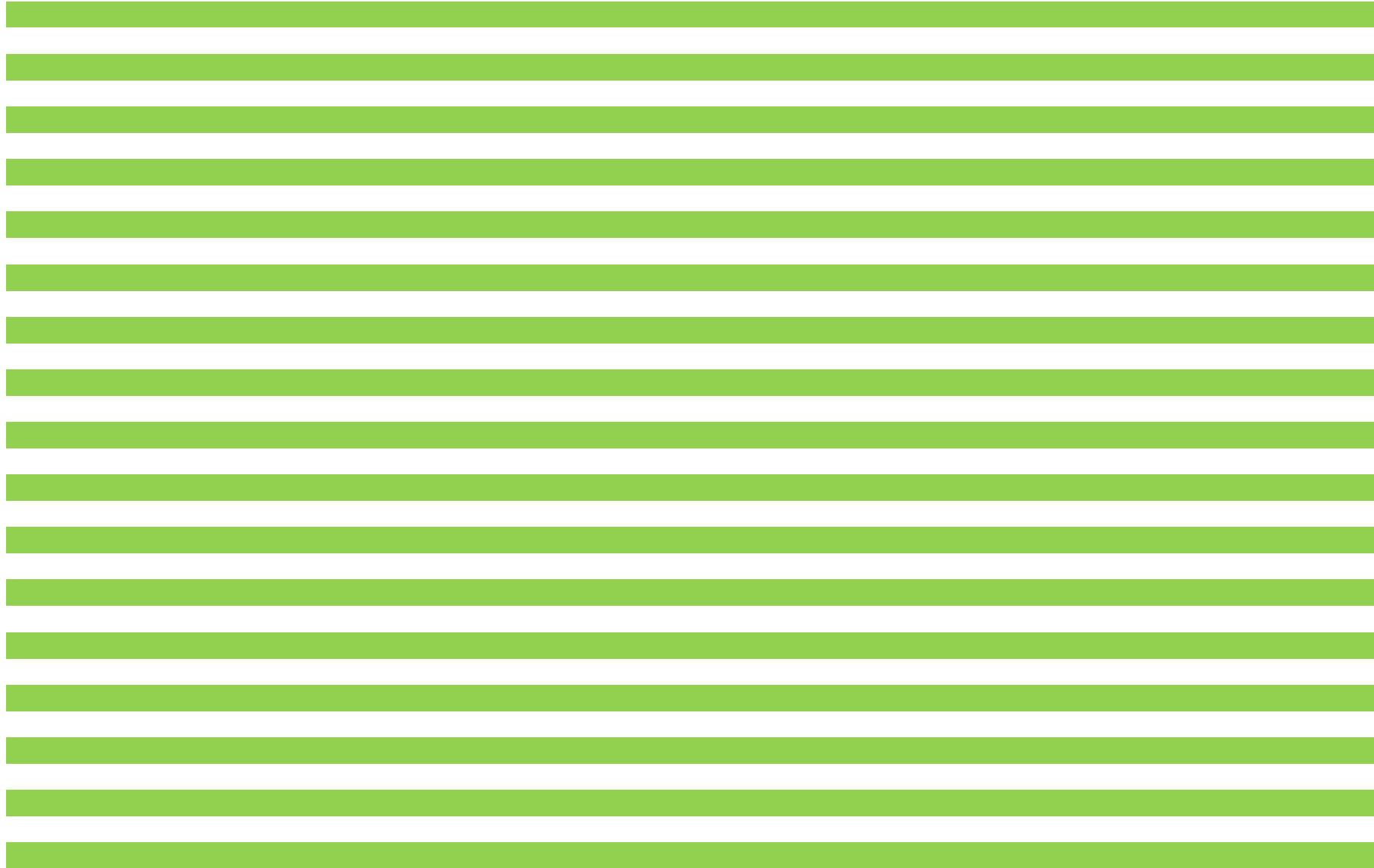
TAB]

TAB]

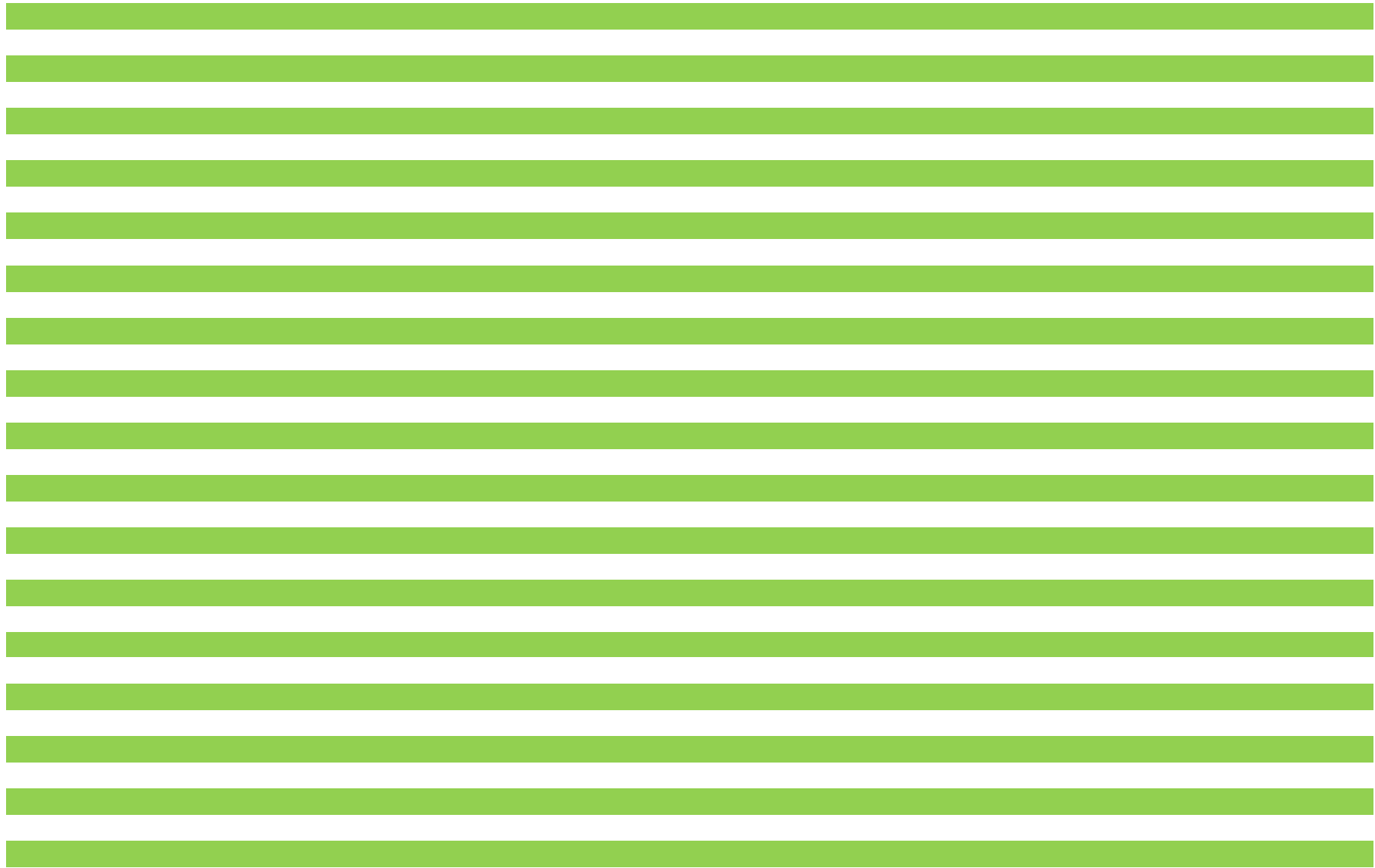
TAB]

TAB]

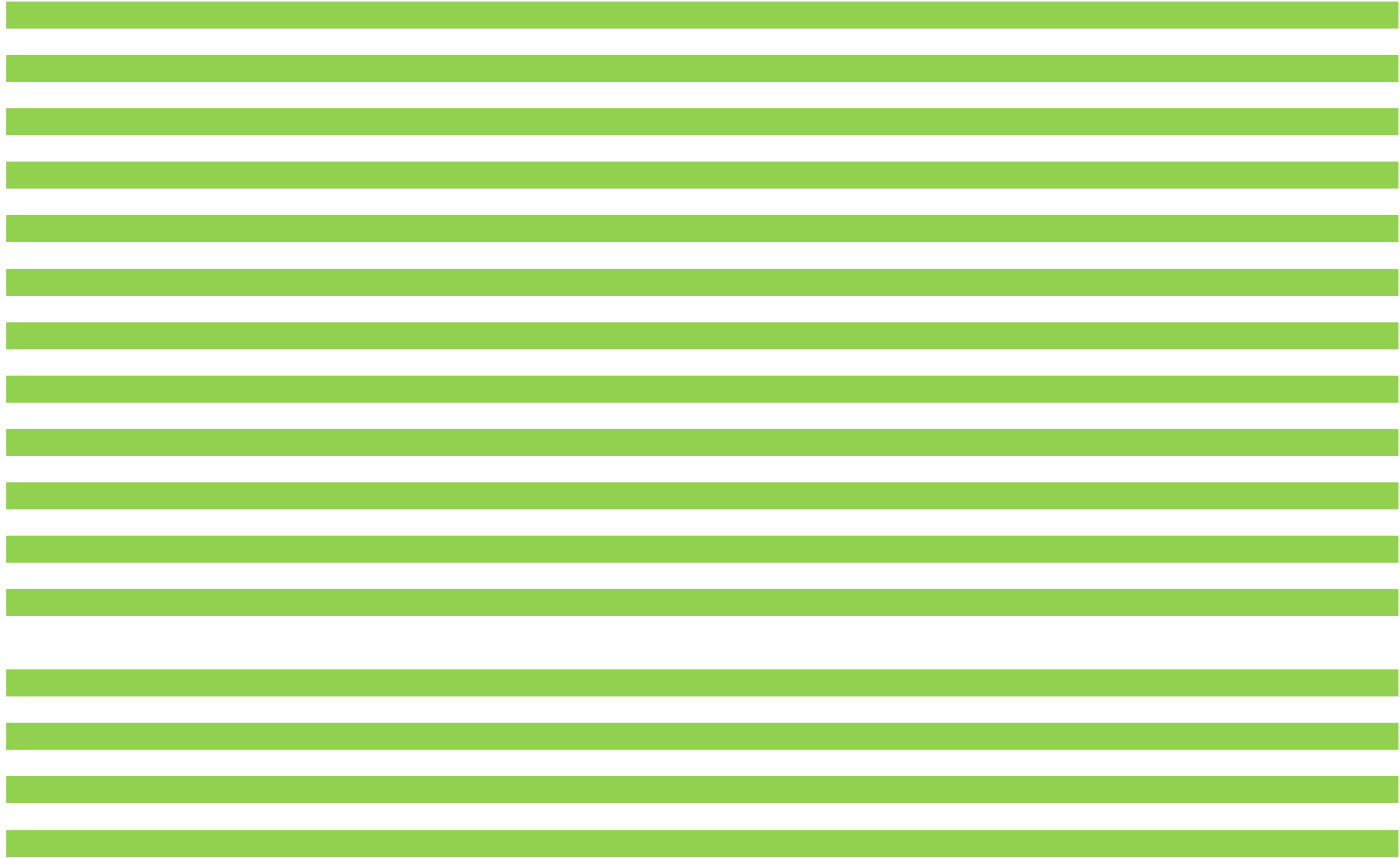
TAB]



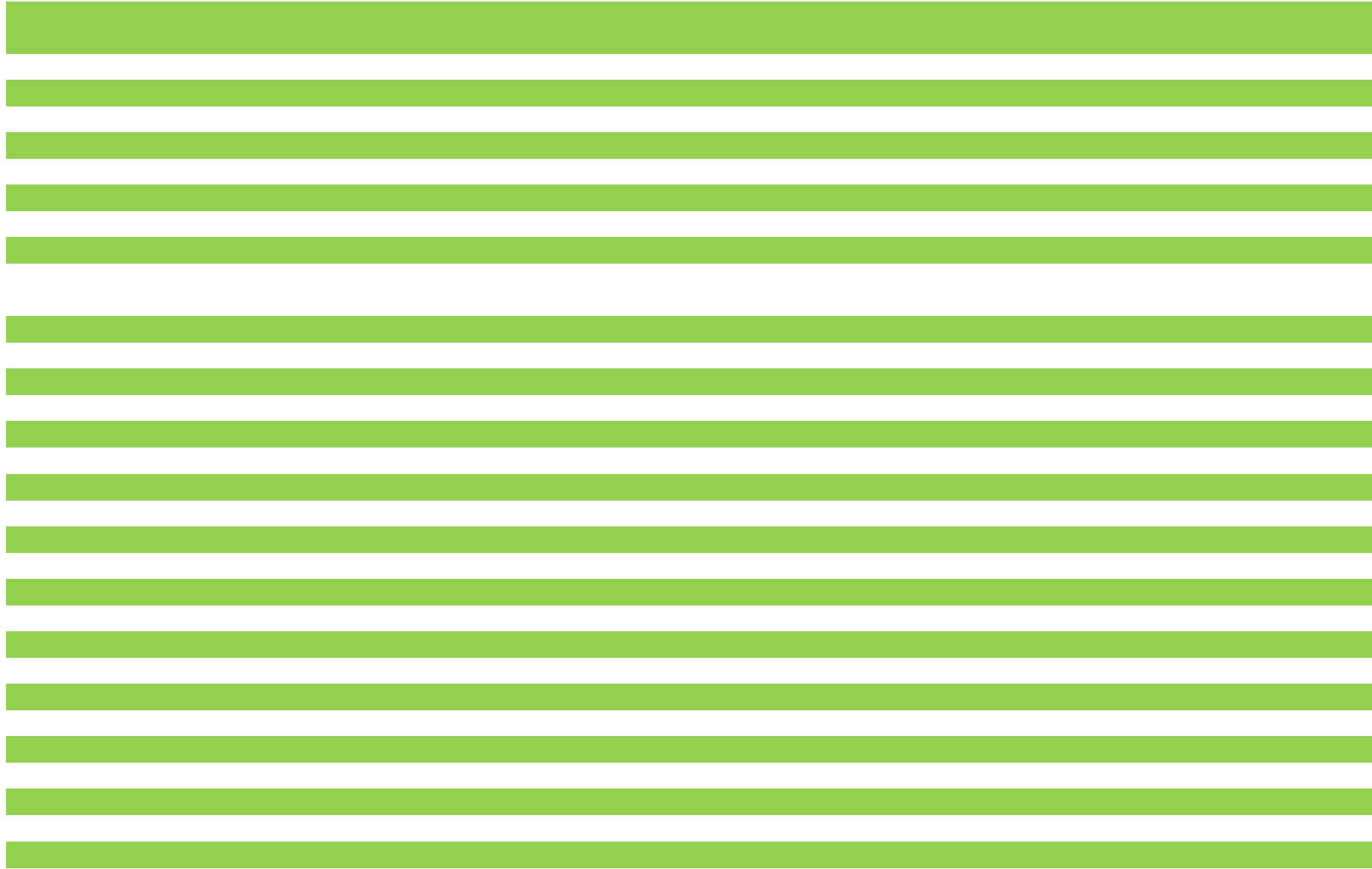
TAB]



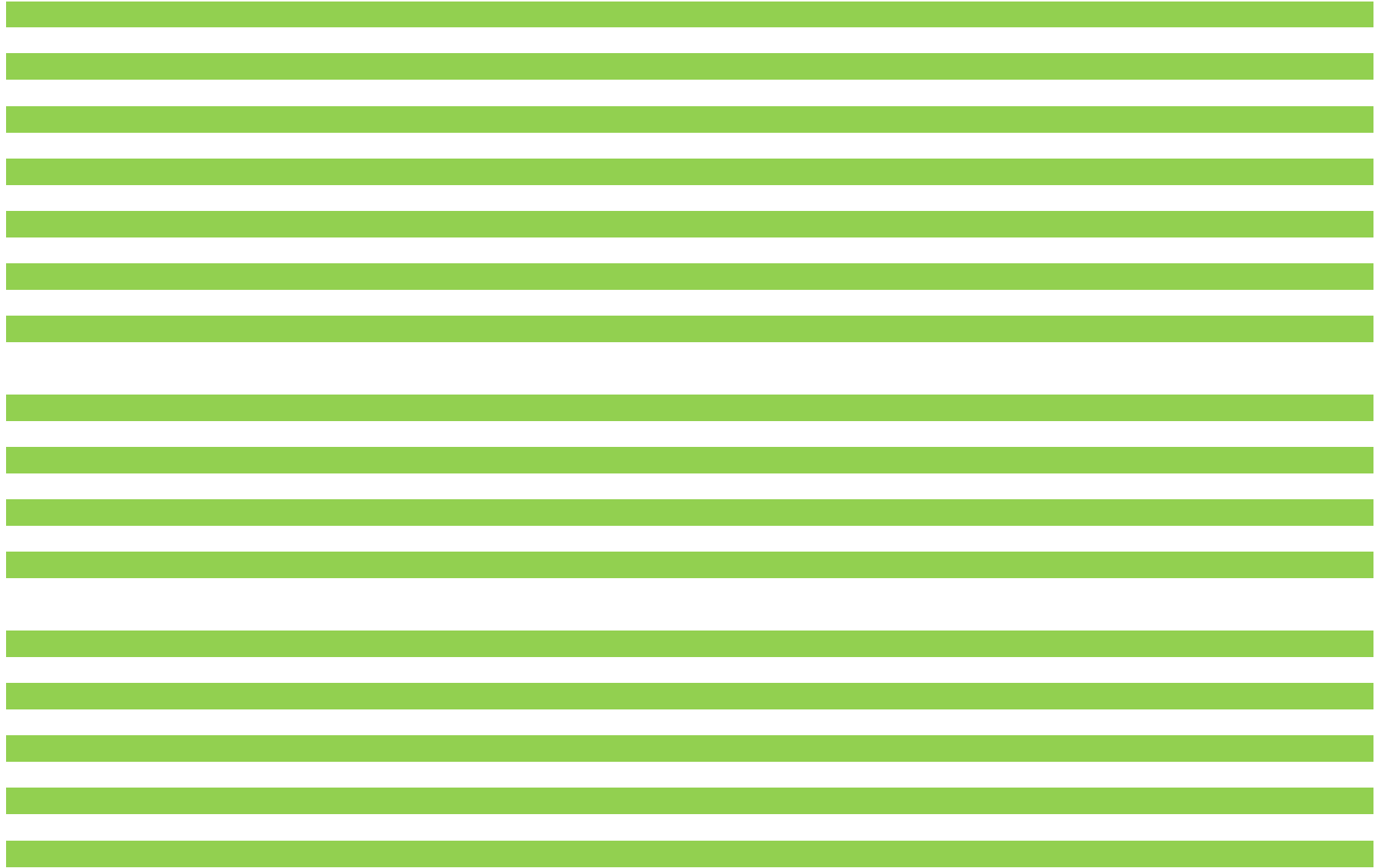
TAB]



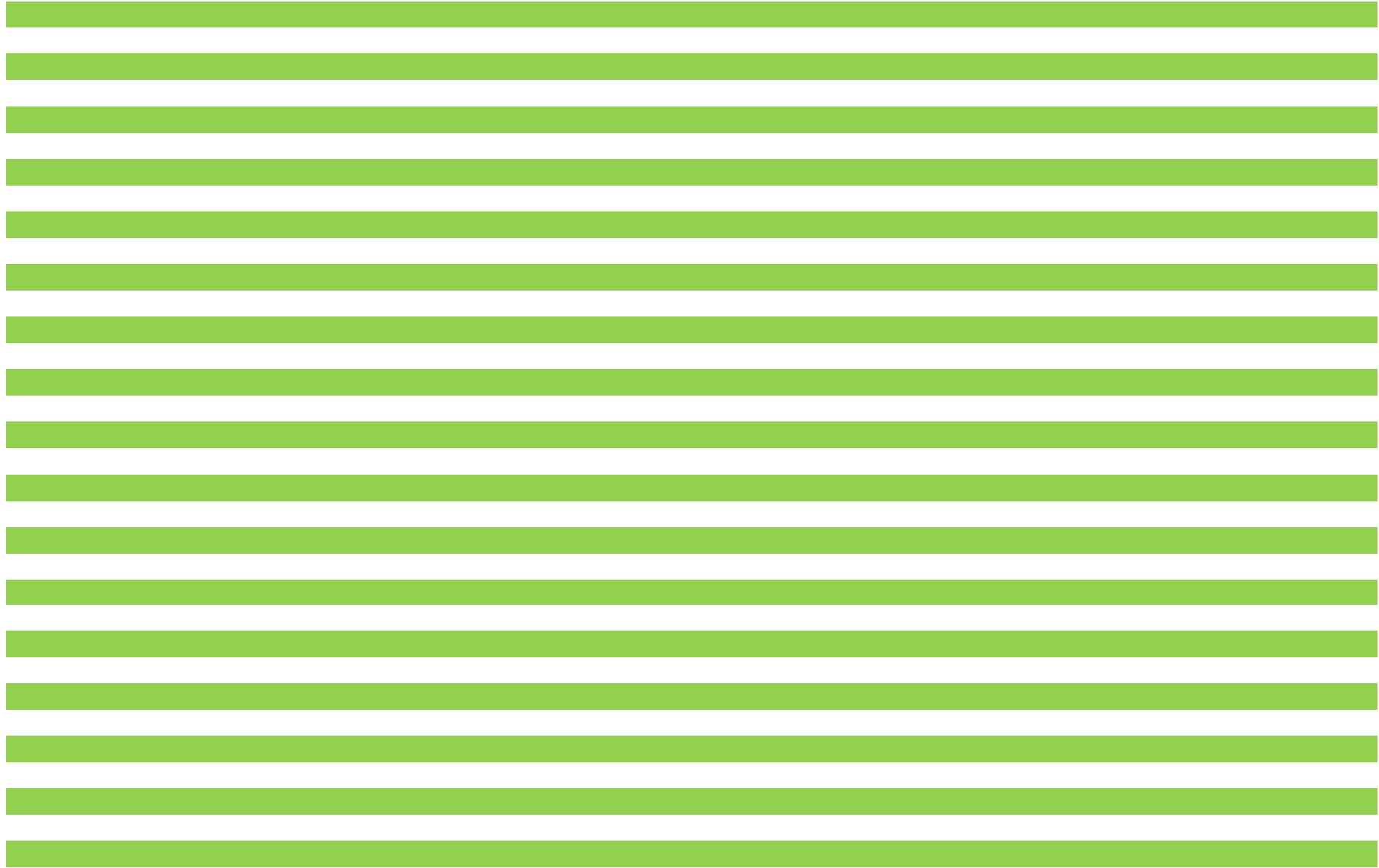
TAB]



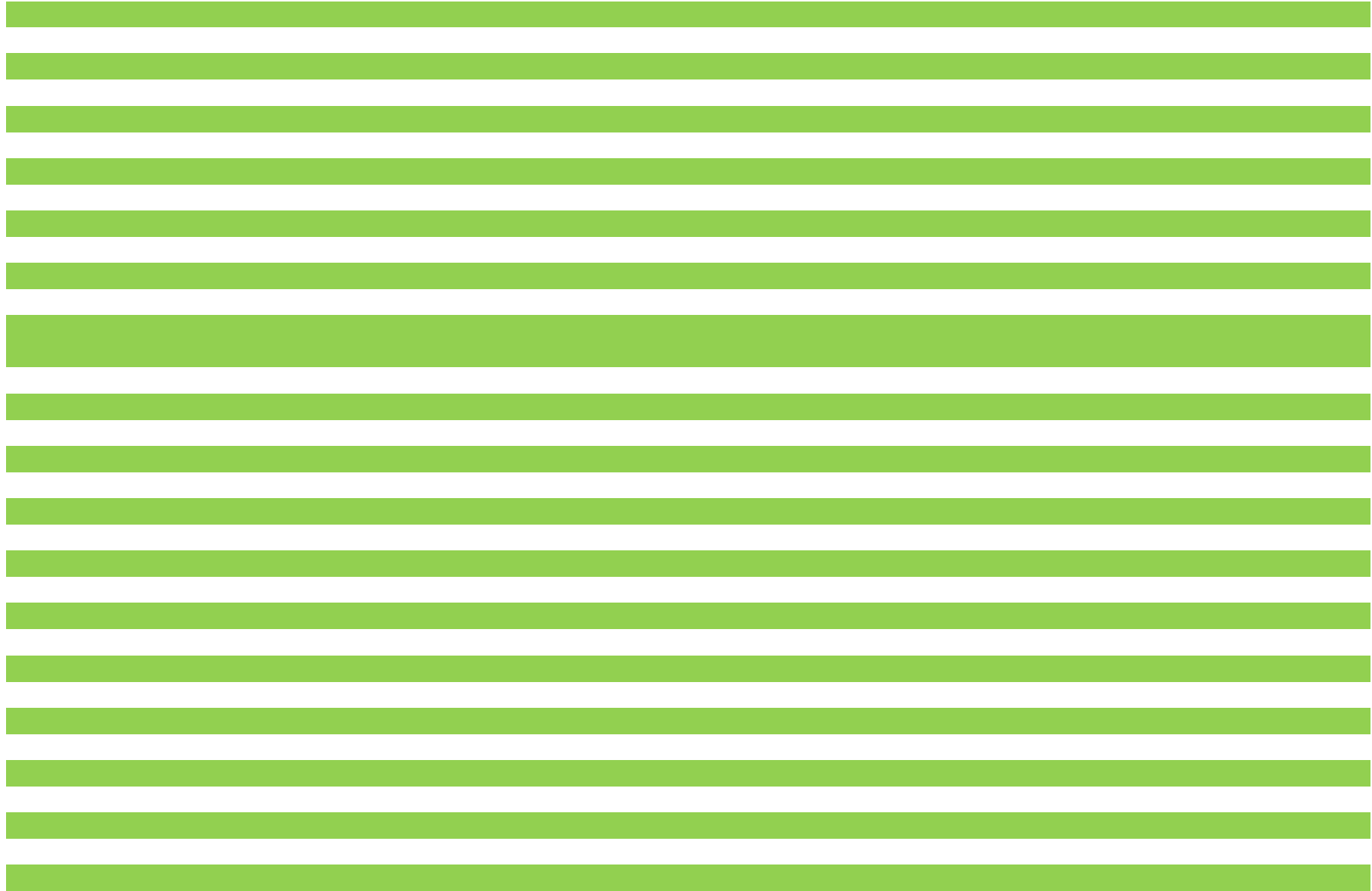
TAB]



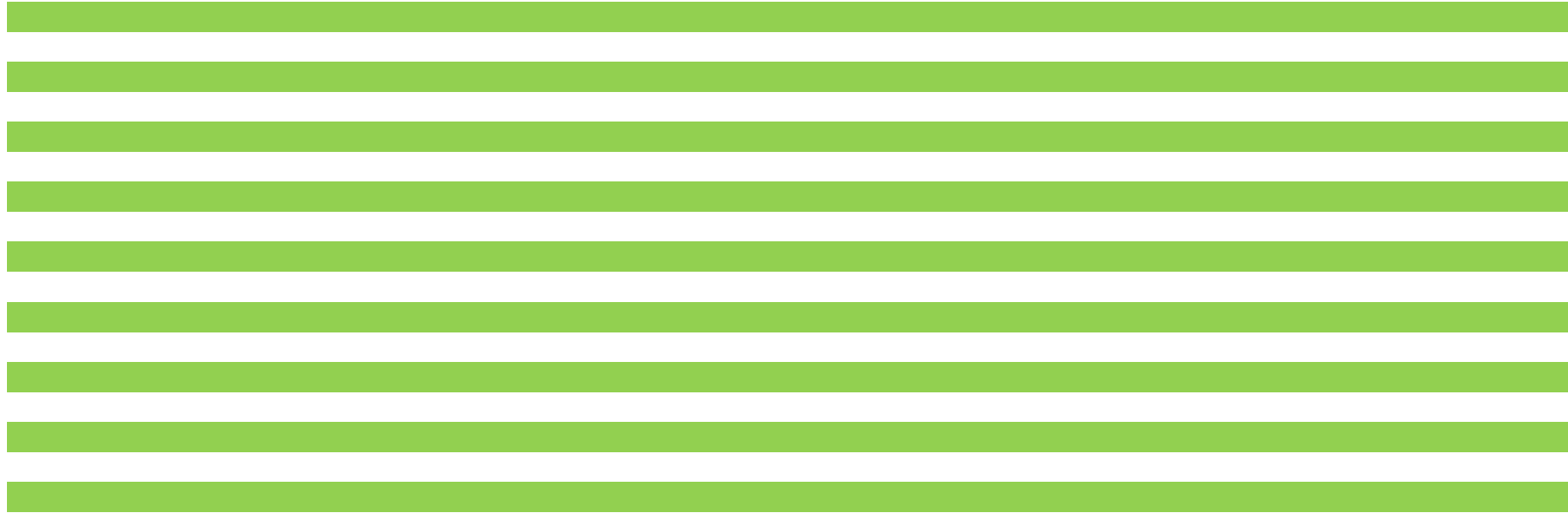
TAB]



TAB]



TAB]



TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

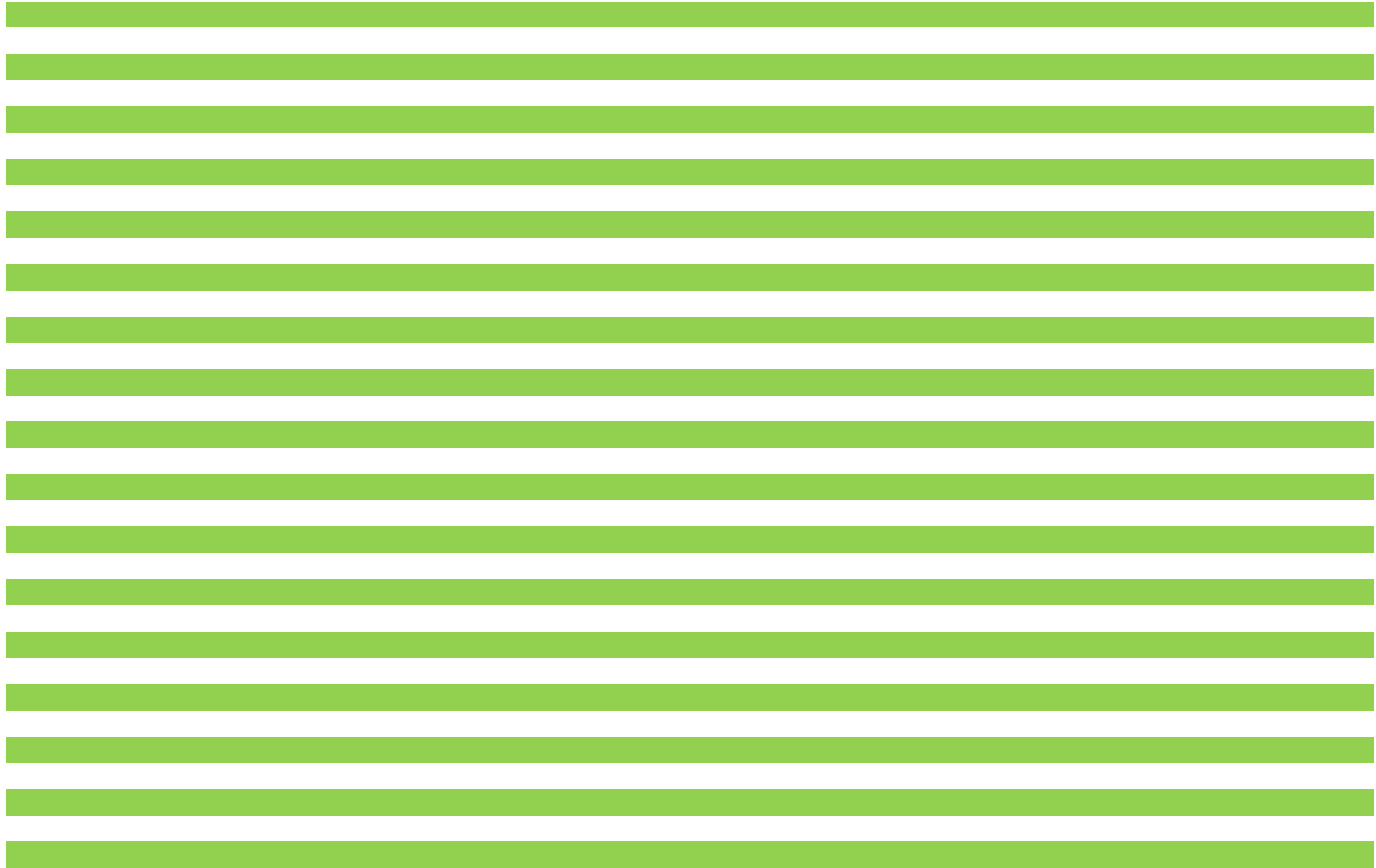
TAB]

TAB]

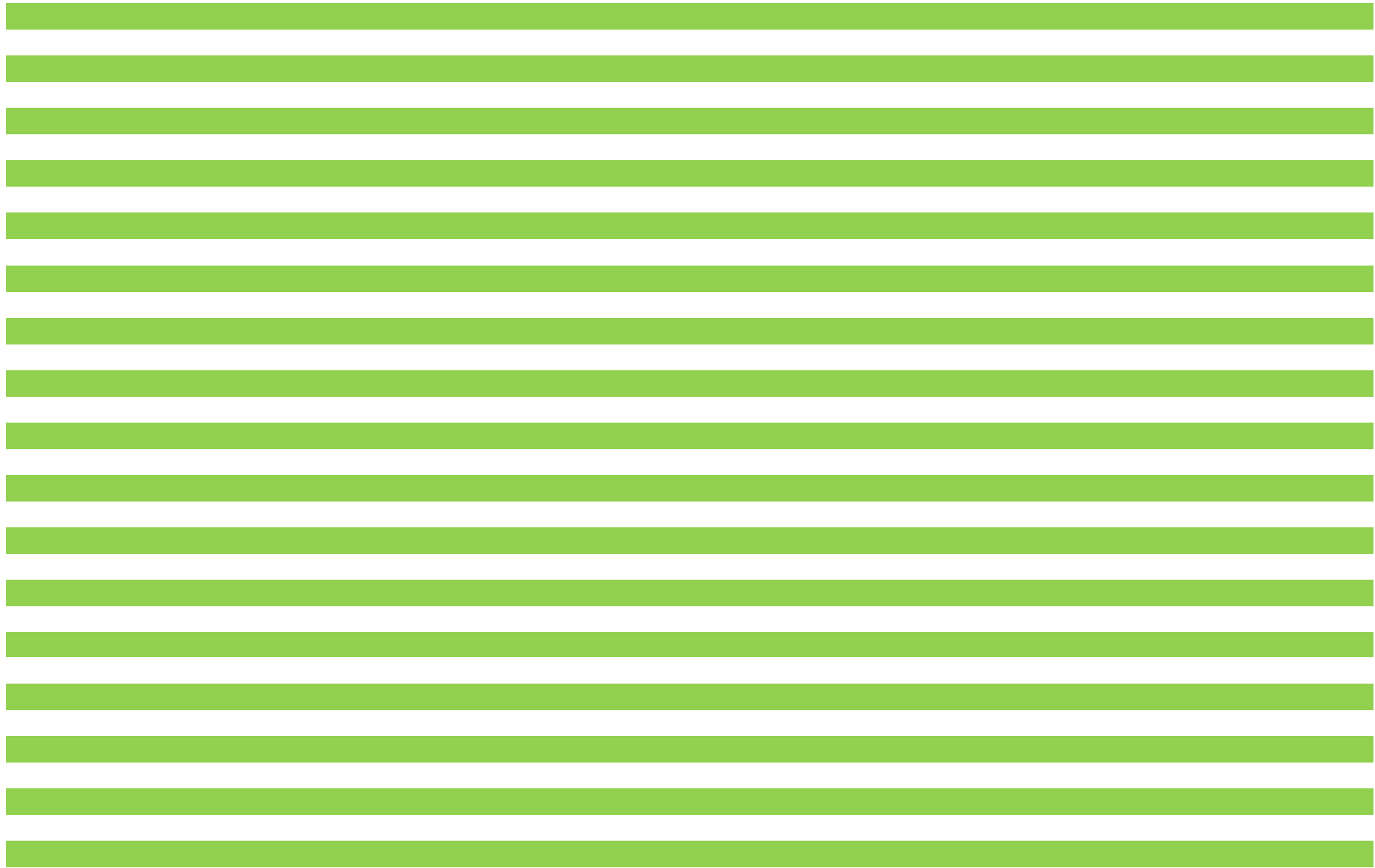
TAB]

TAB]

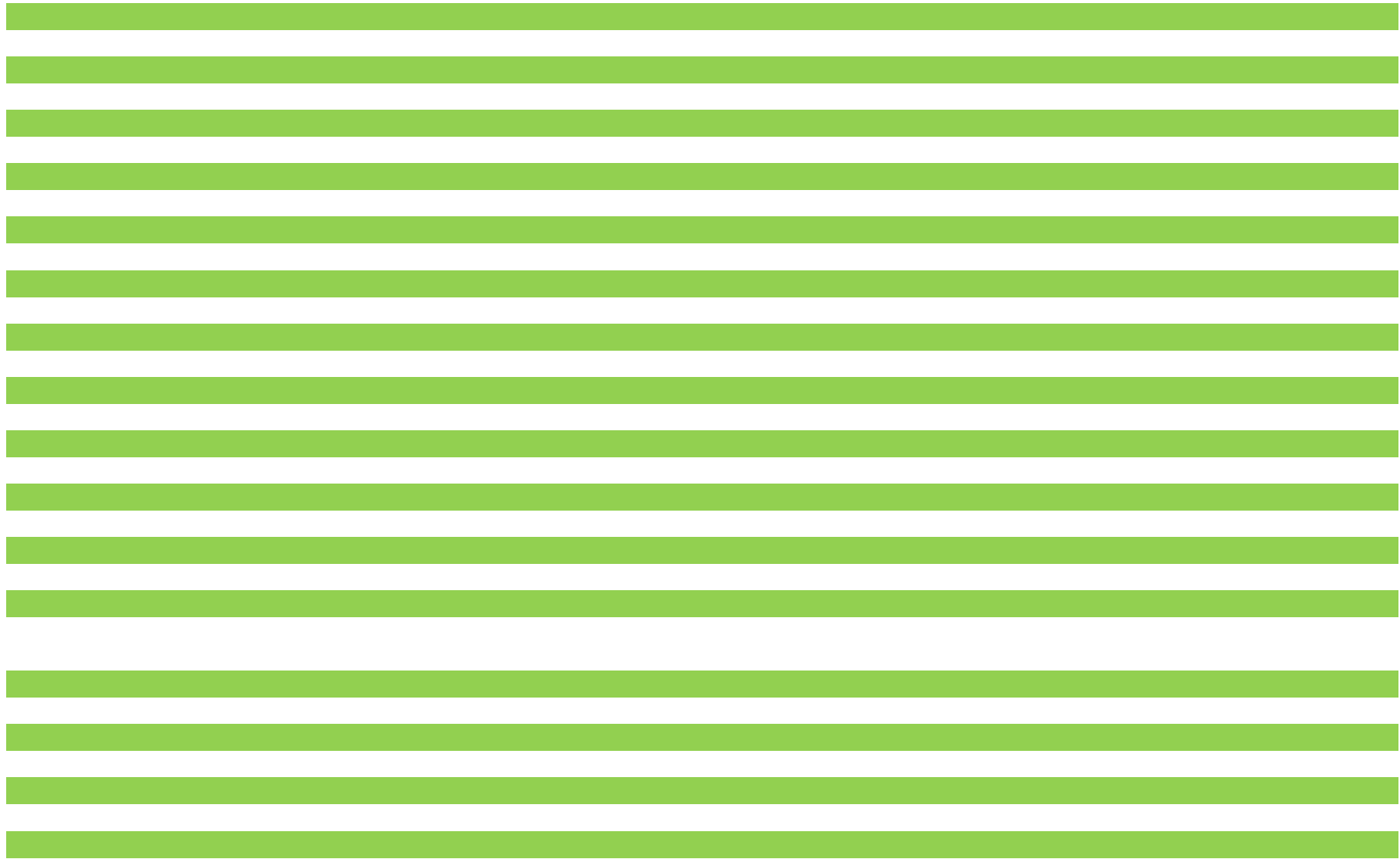
TAB]



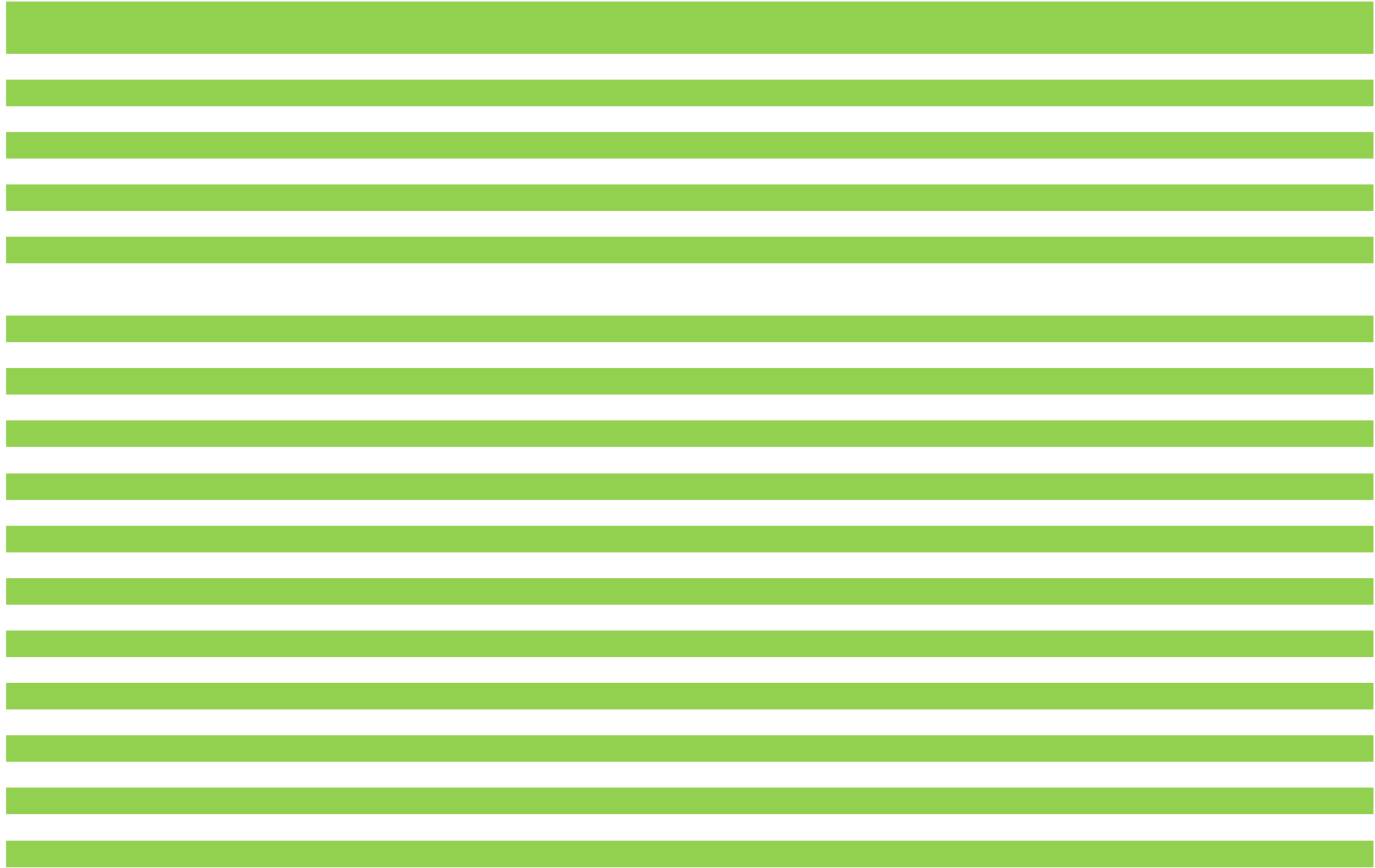
TAB]



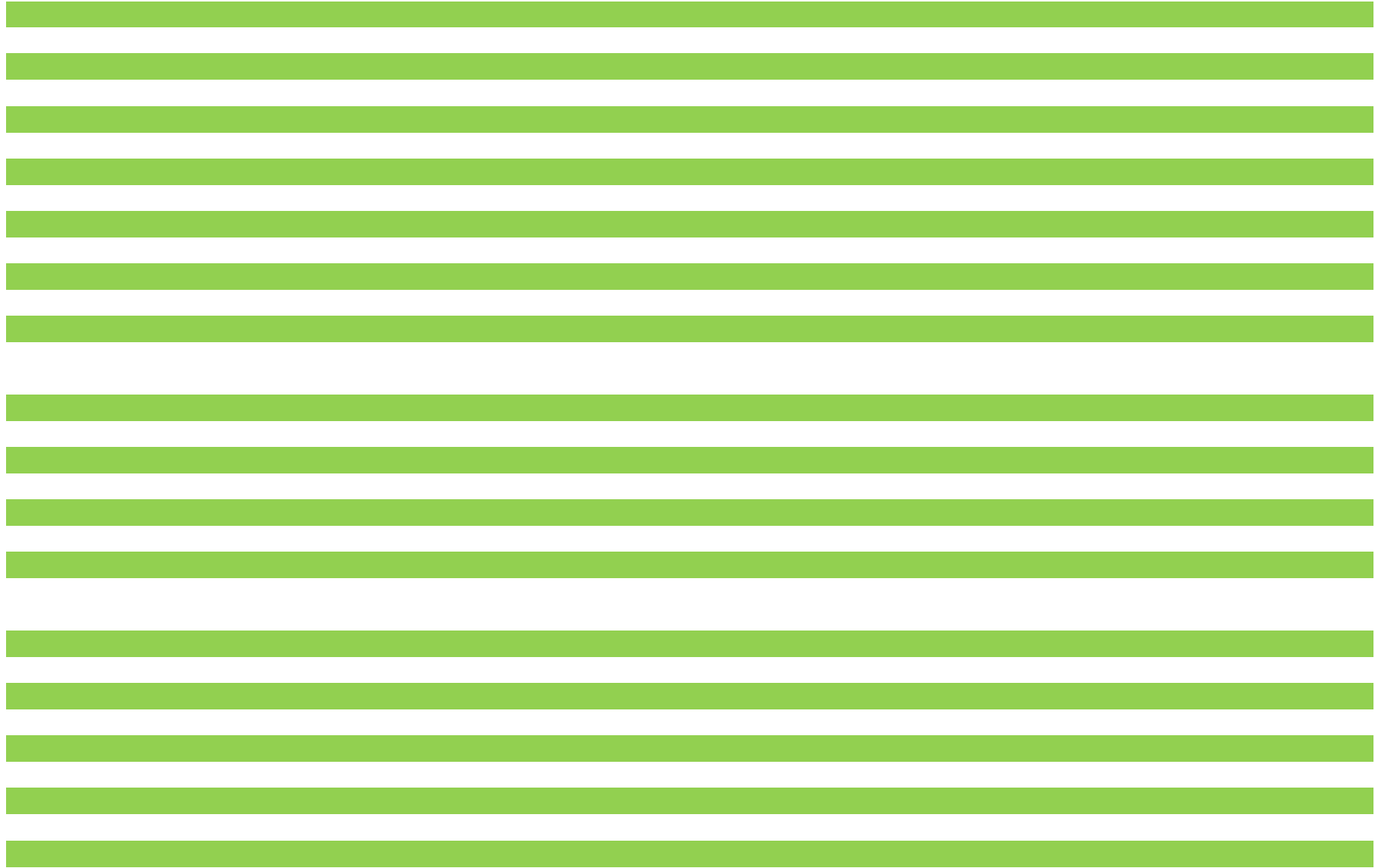
TAB]



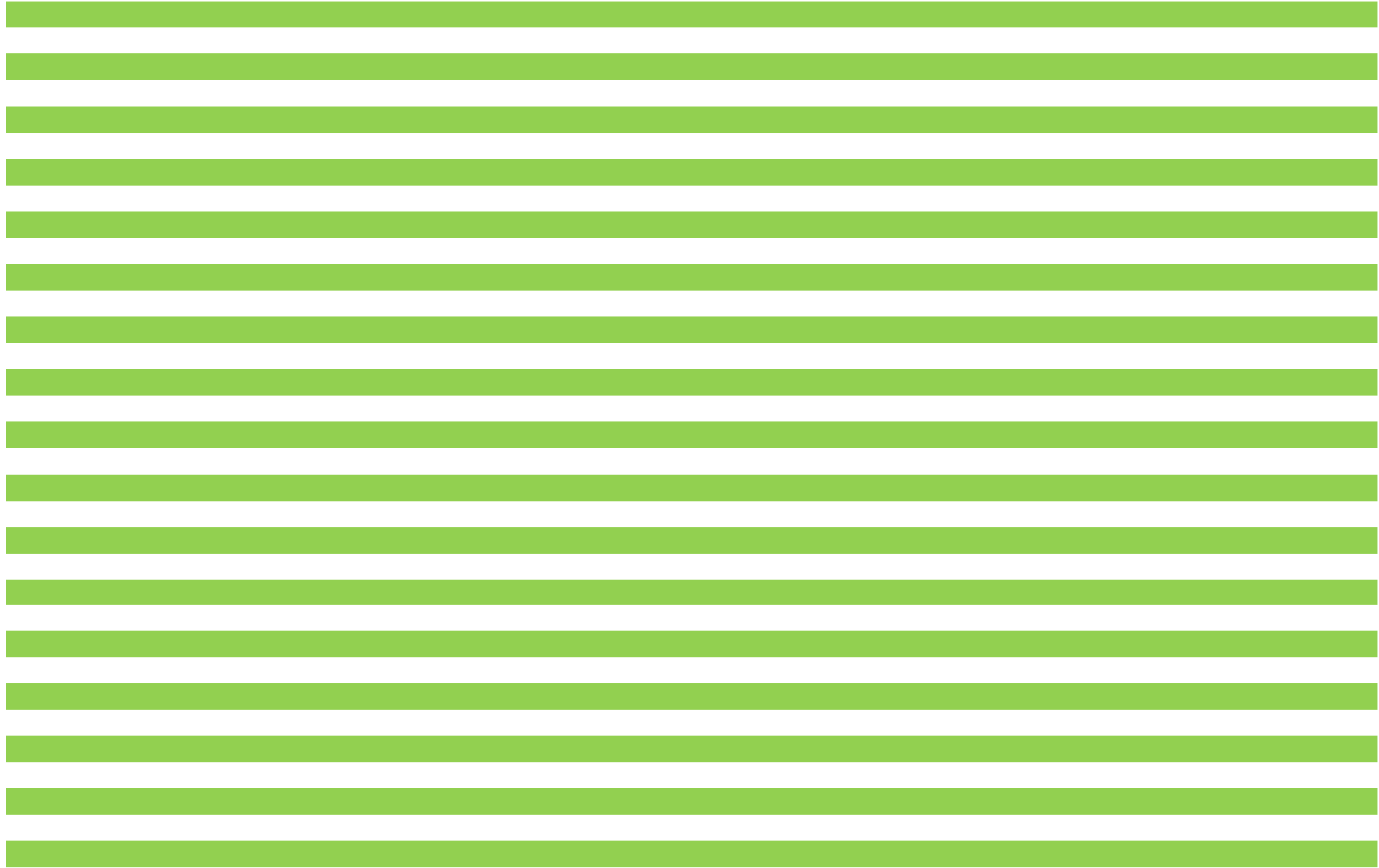
TAB]



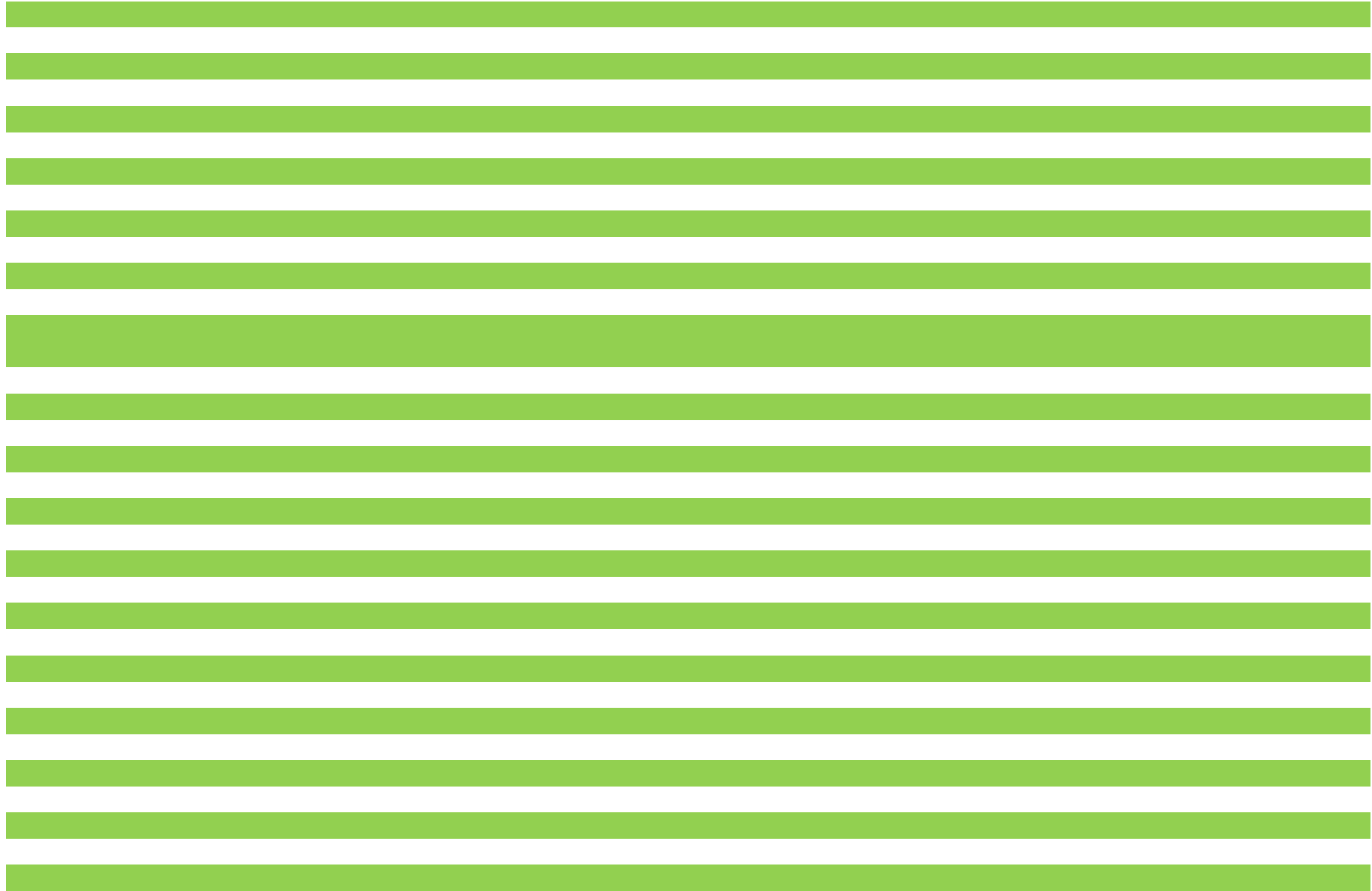
TAB]



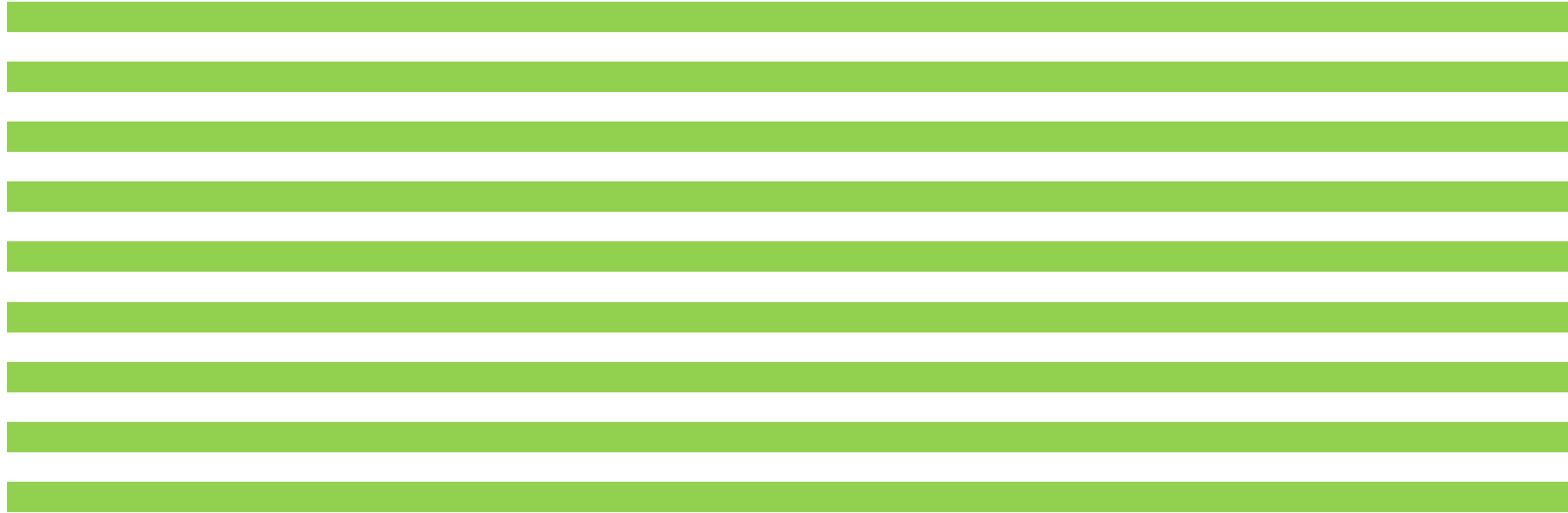
TAB]



TAB]



TAB]



TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

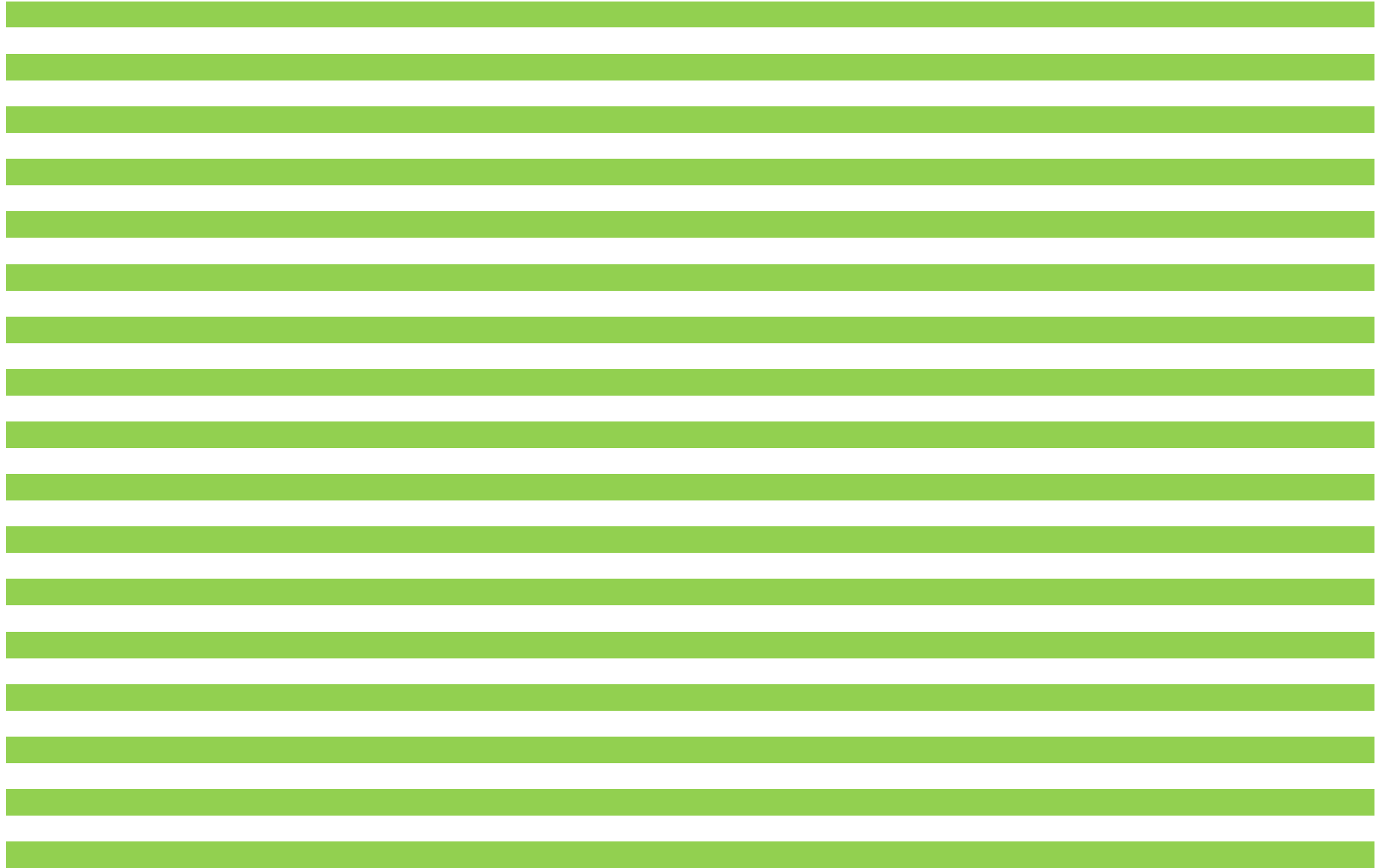
TAB]

TAB]

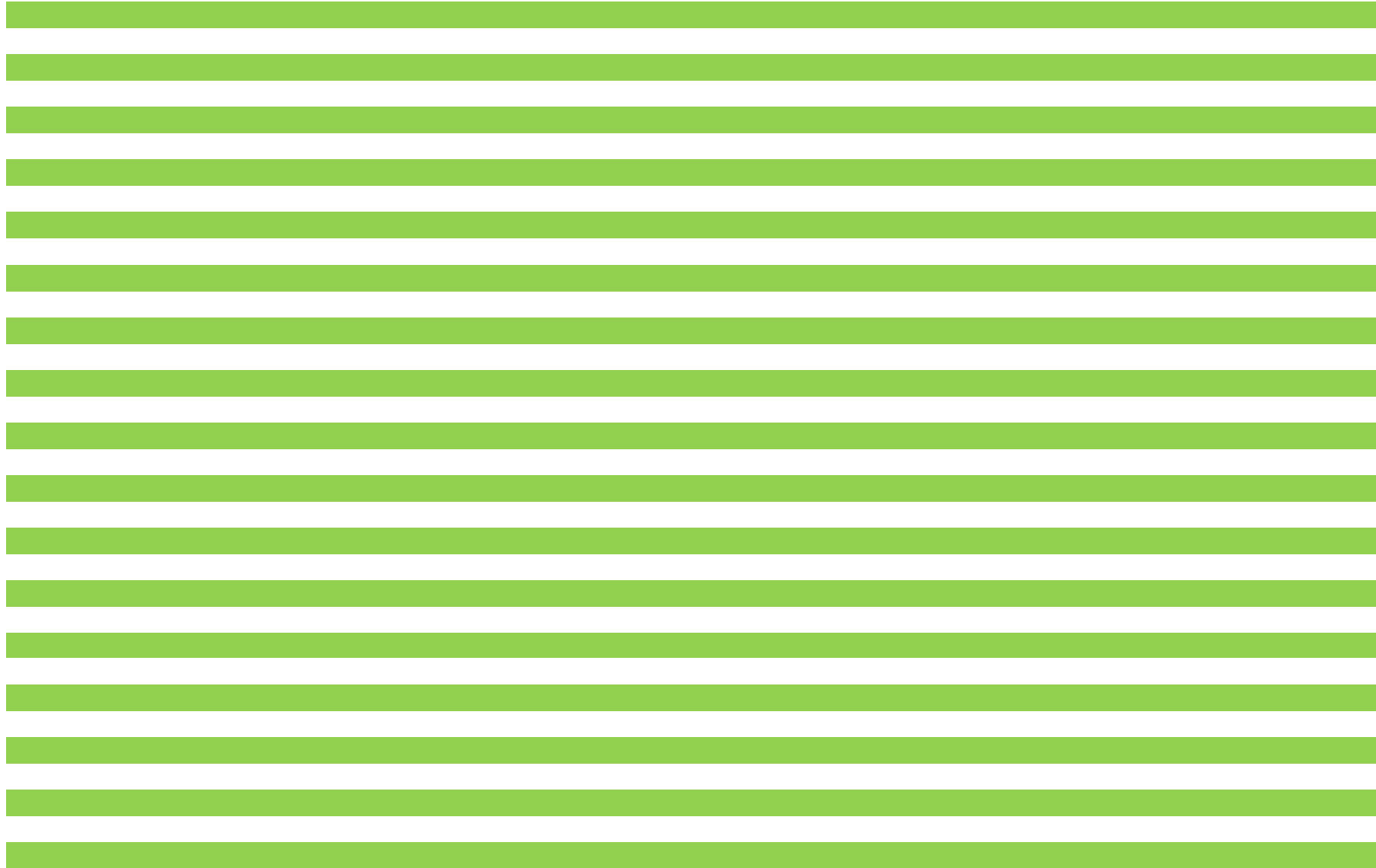
TAB]

TAB]

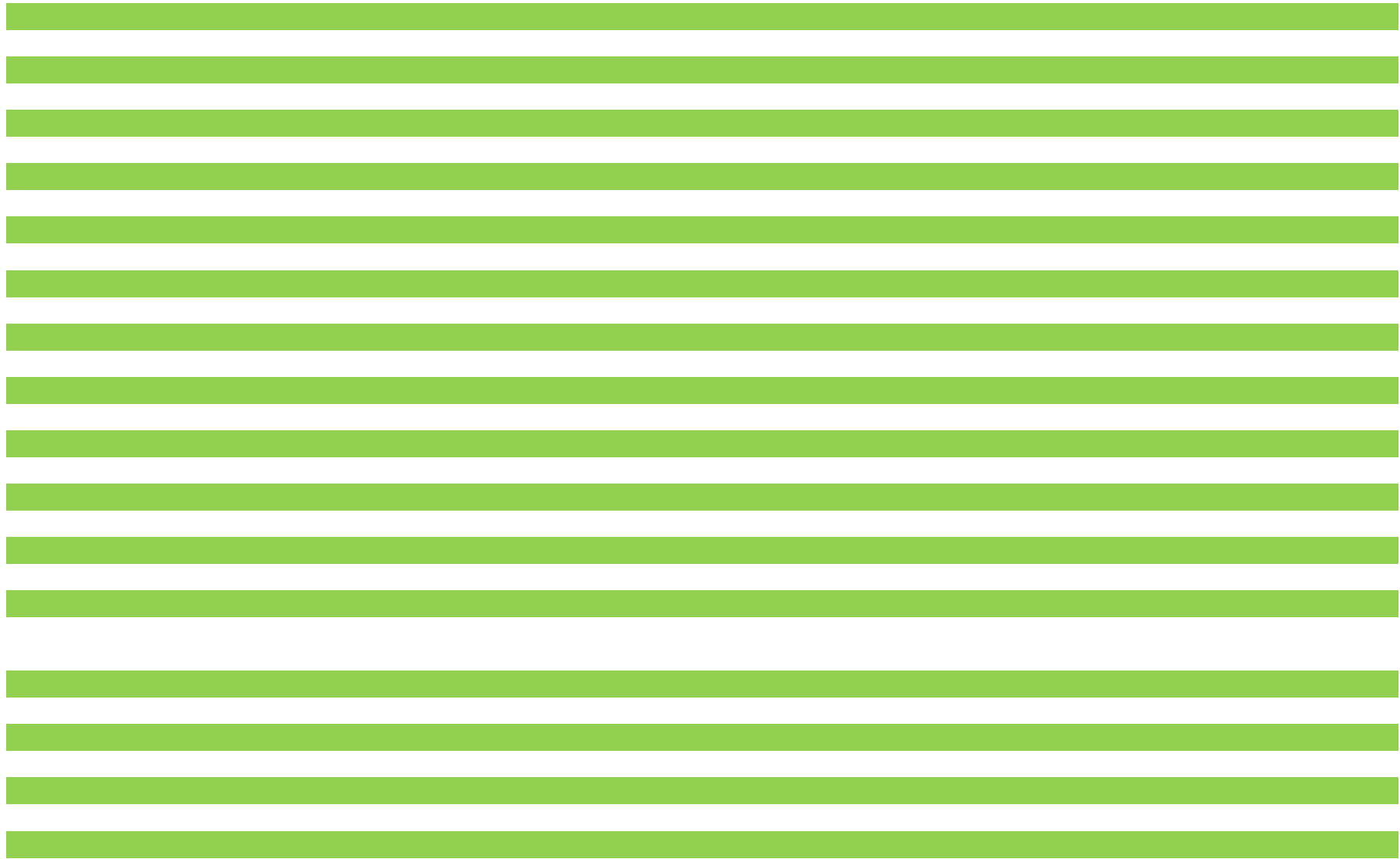
TAB]



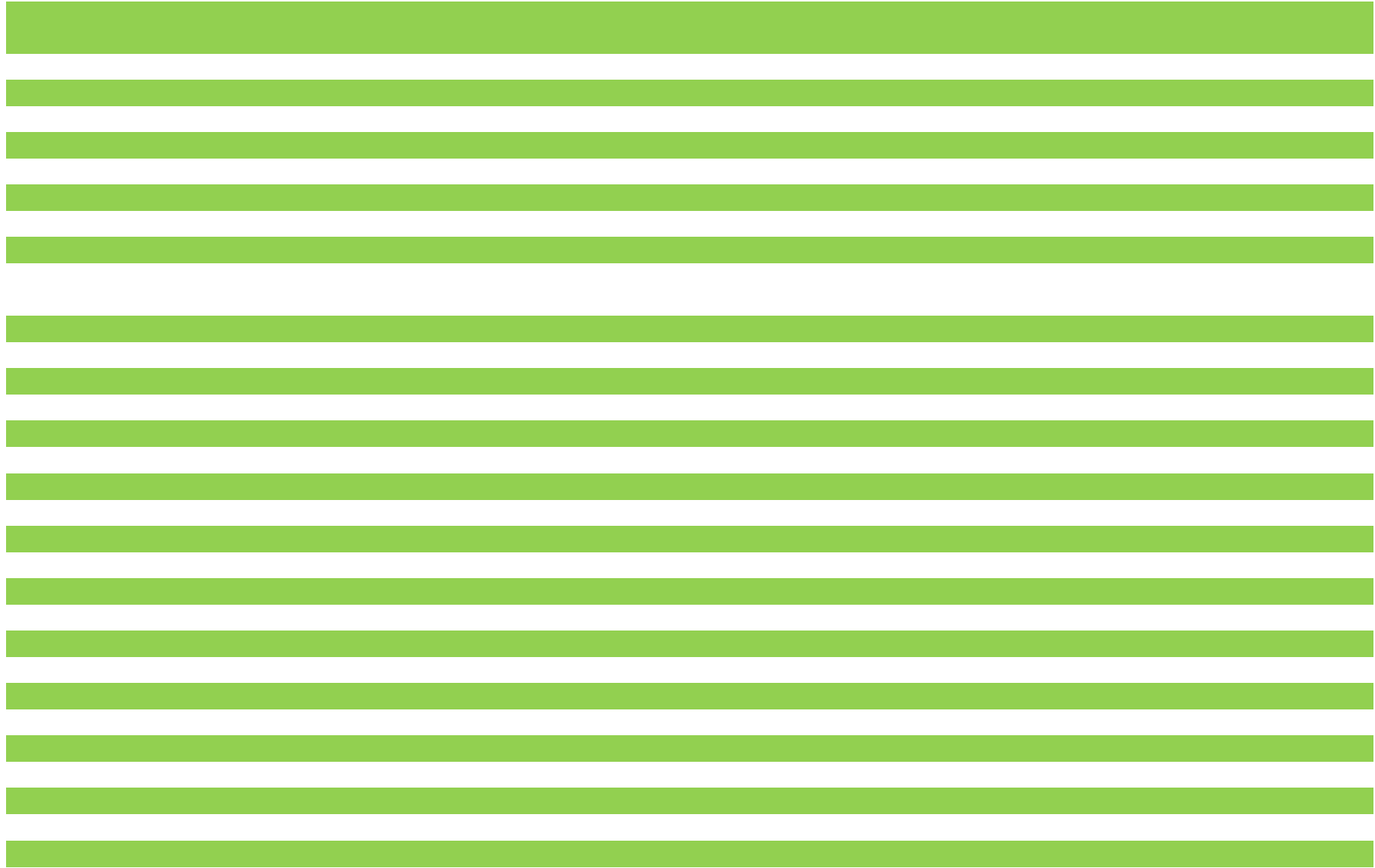
TAB]



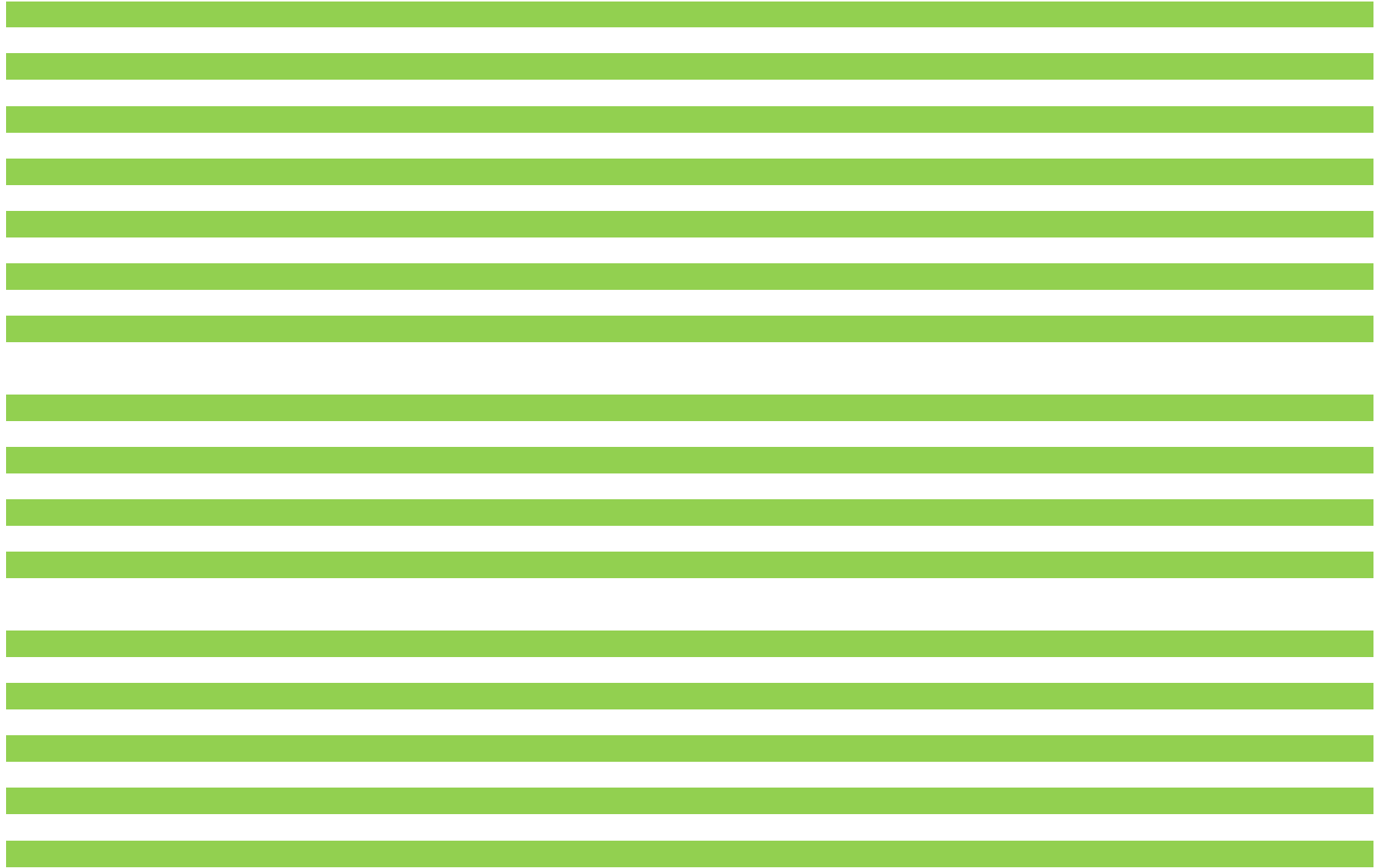
TAB]



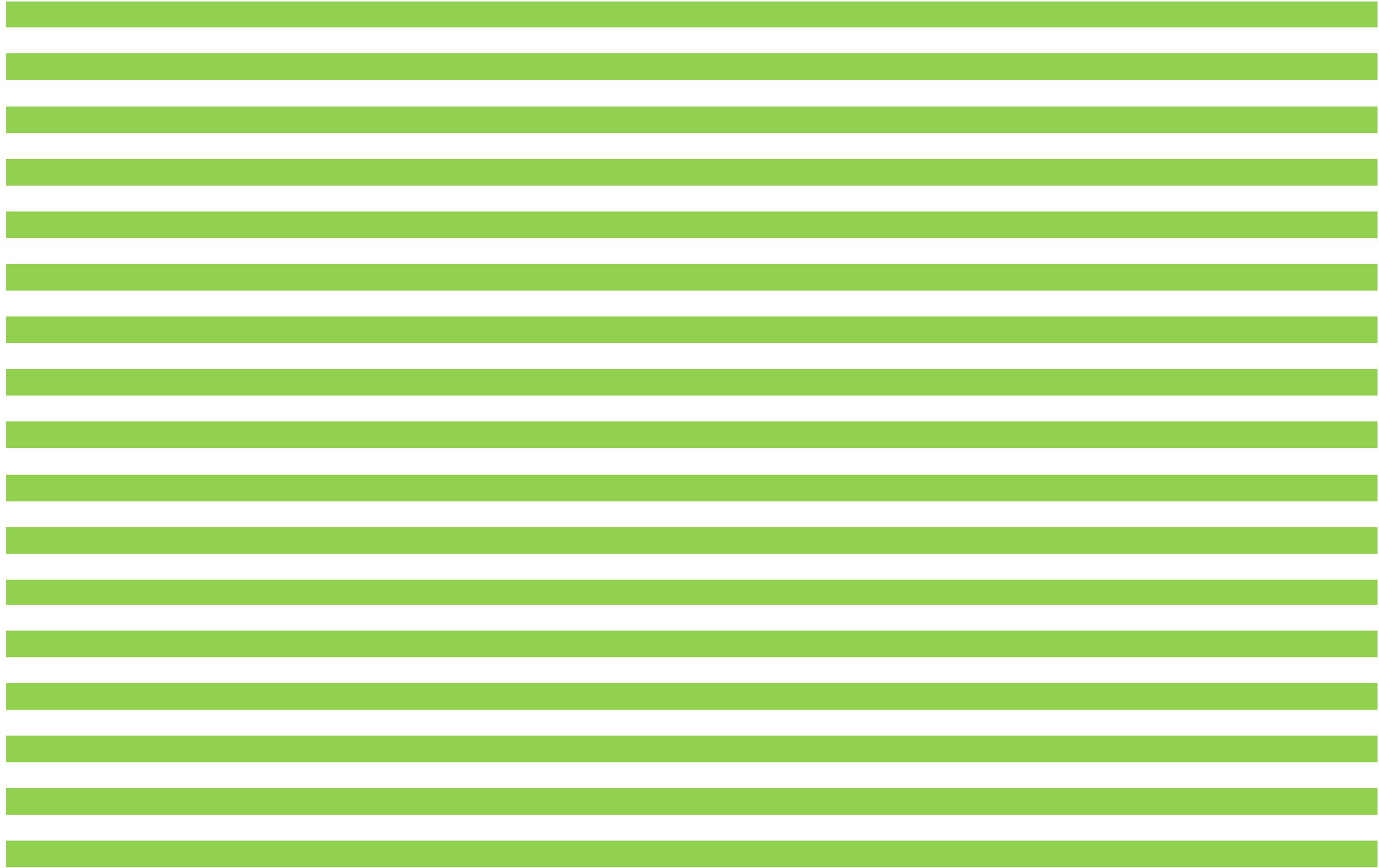
TAB]



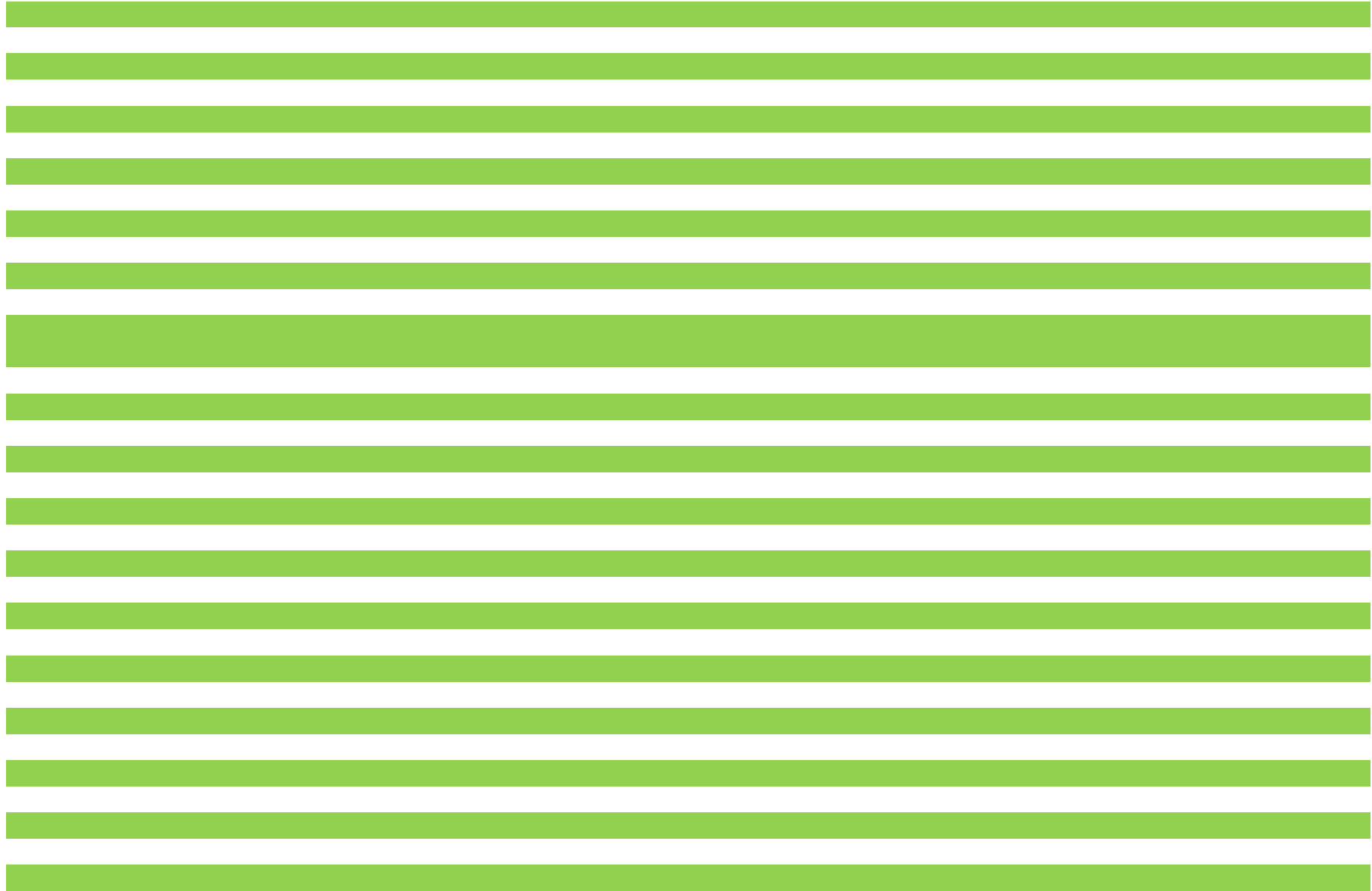
TAB]



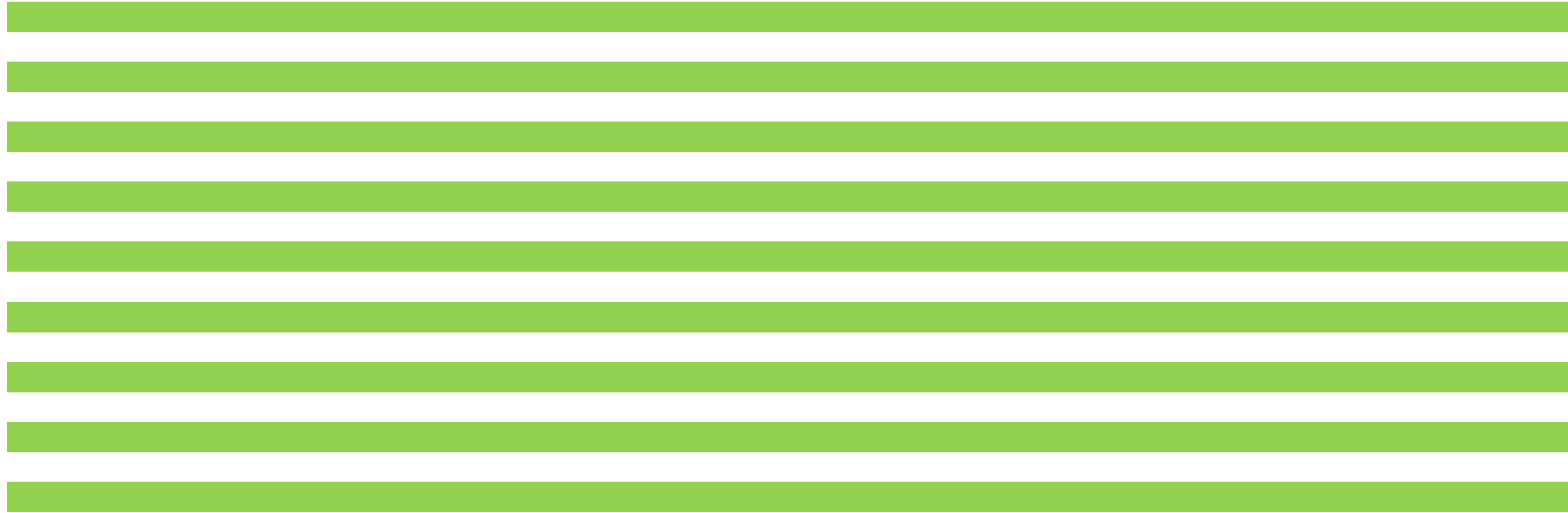
TAB]



TAB]



TAB]



TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

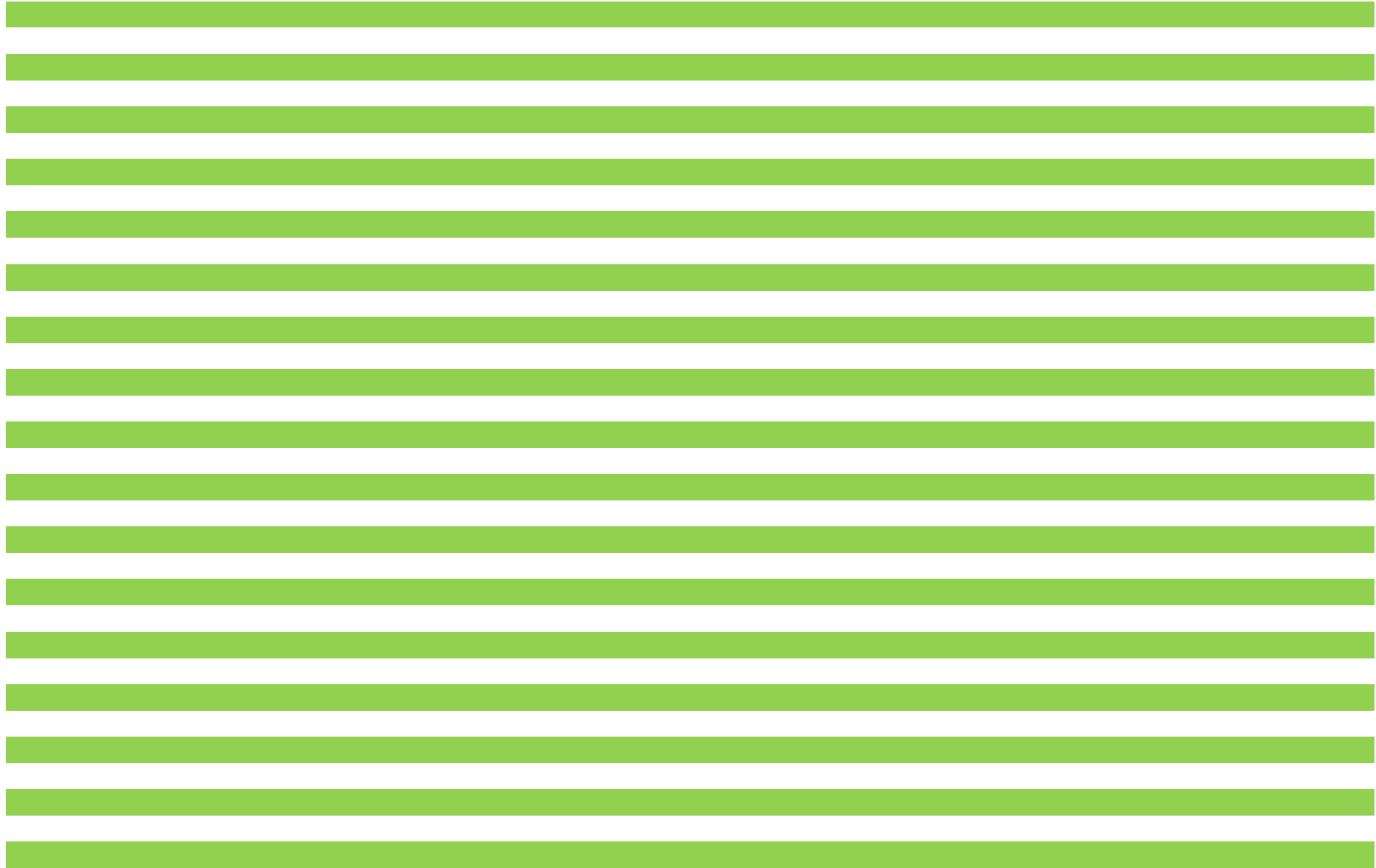
TAB]

TAB]

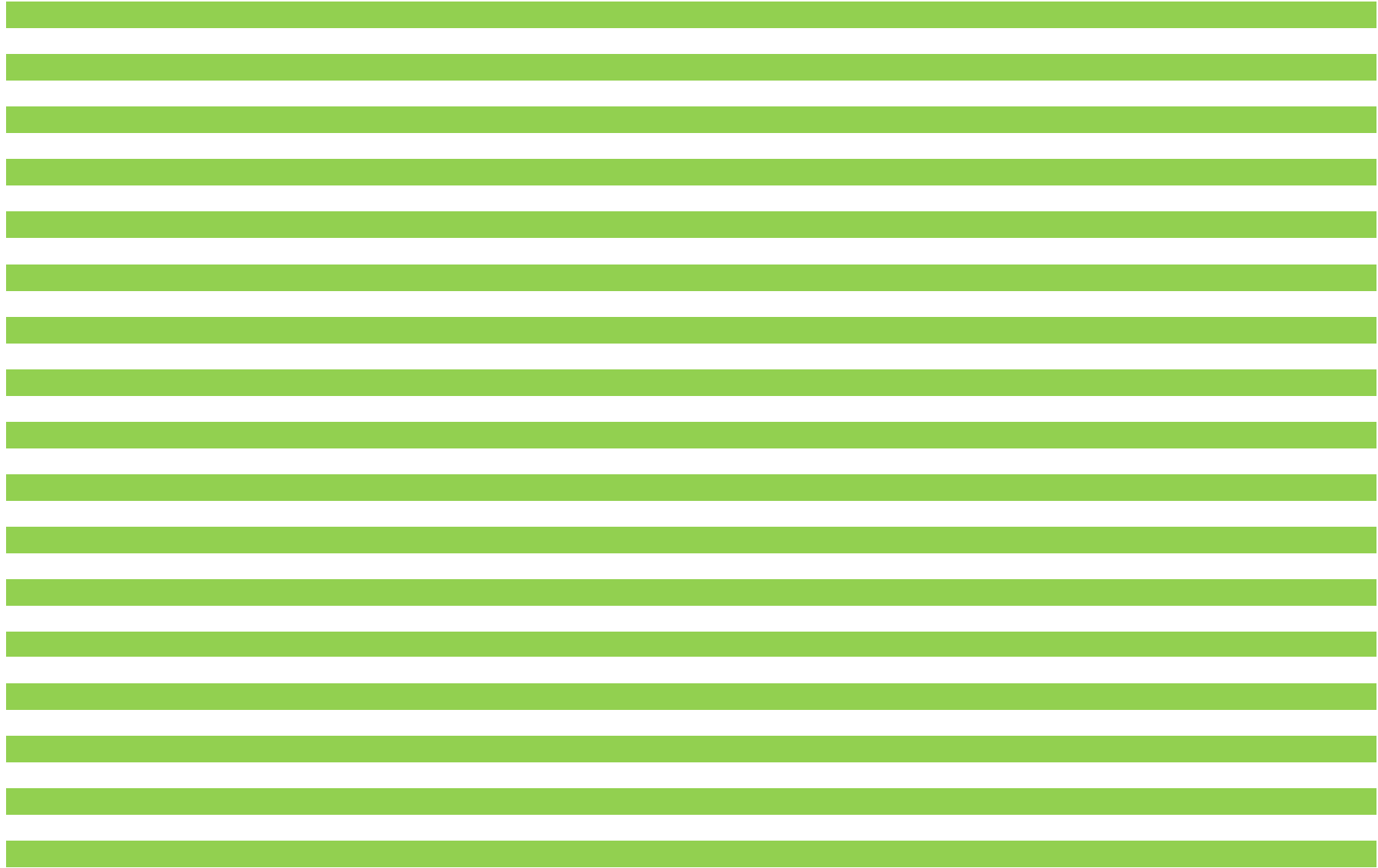
TAB]

TAB]

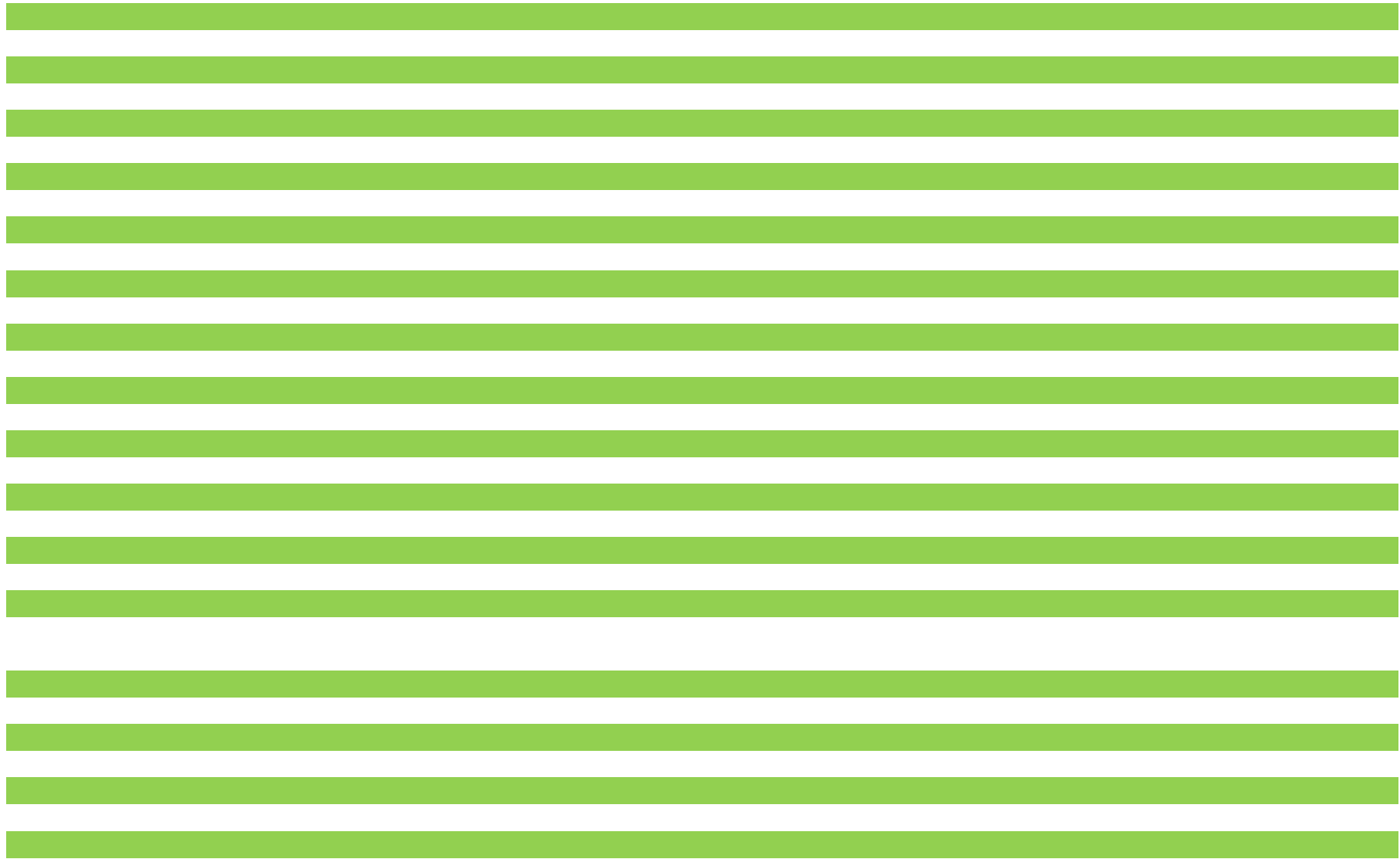
TAB]



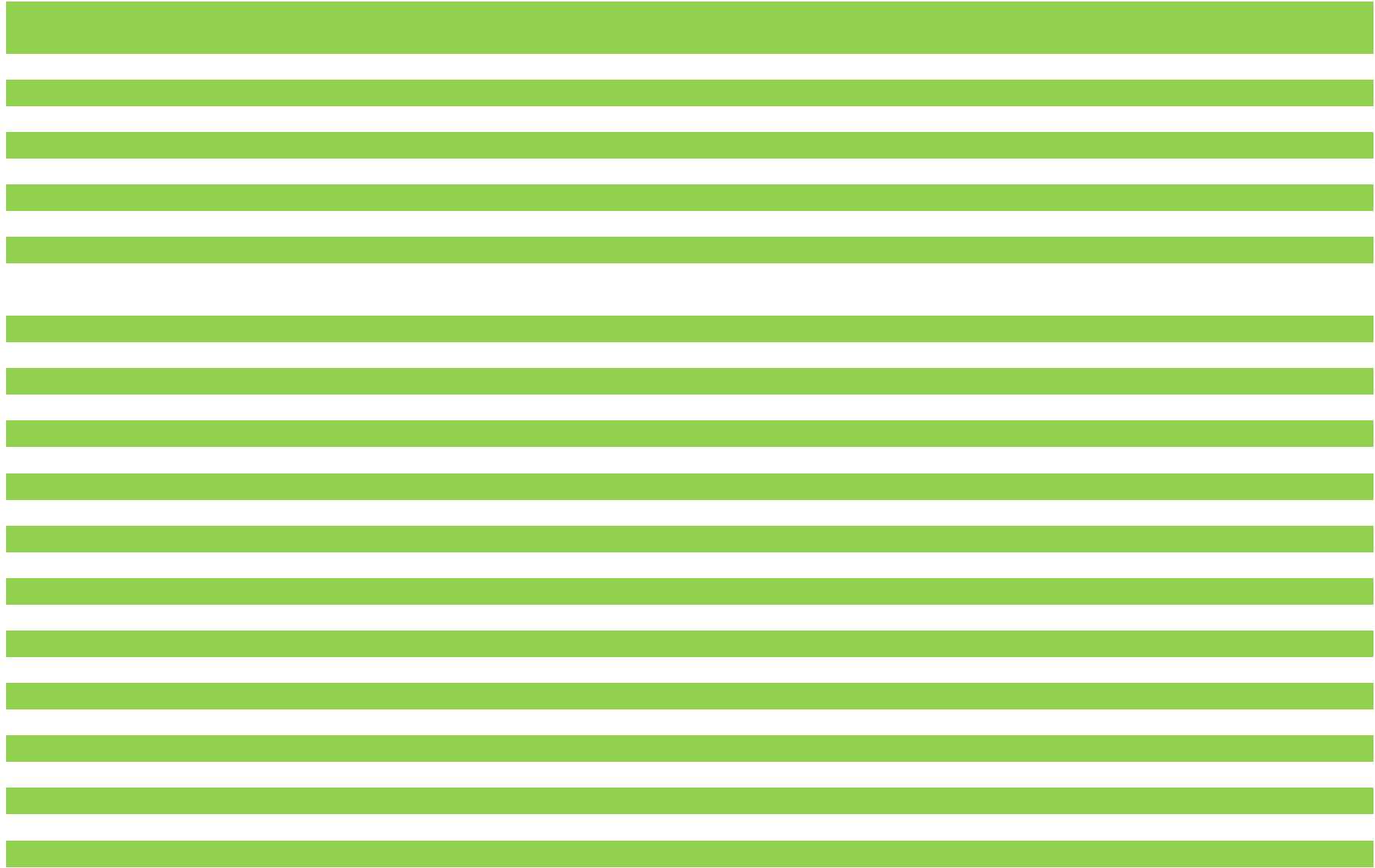
TAB]



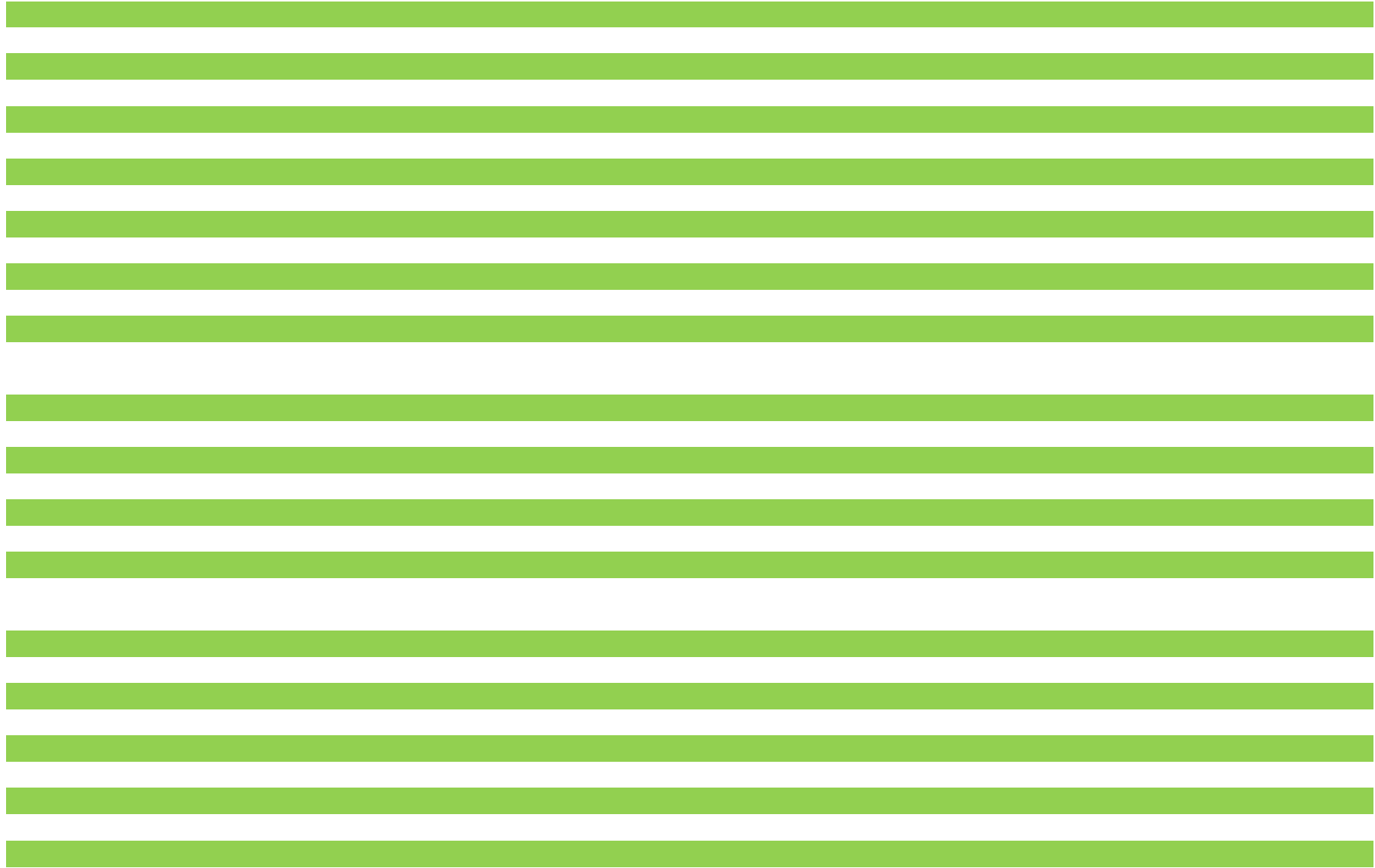
TAB]



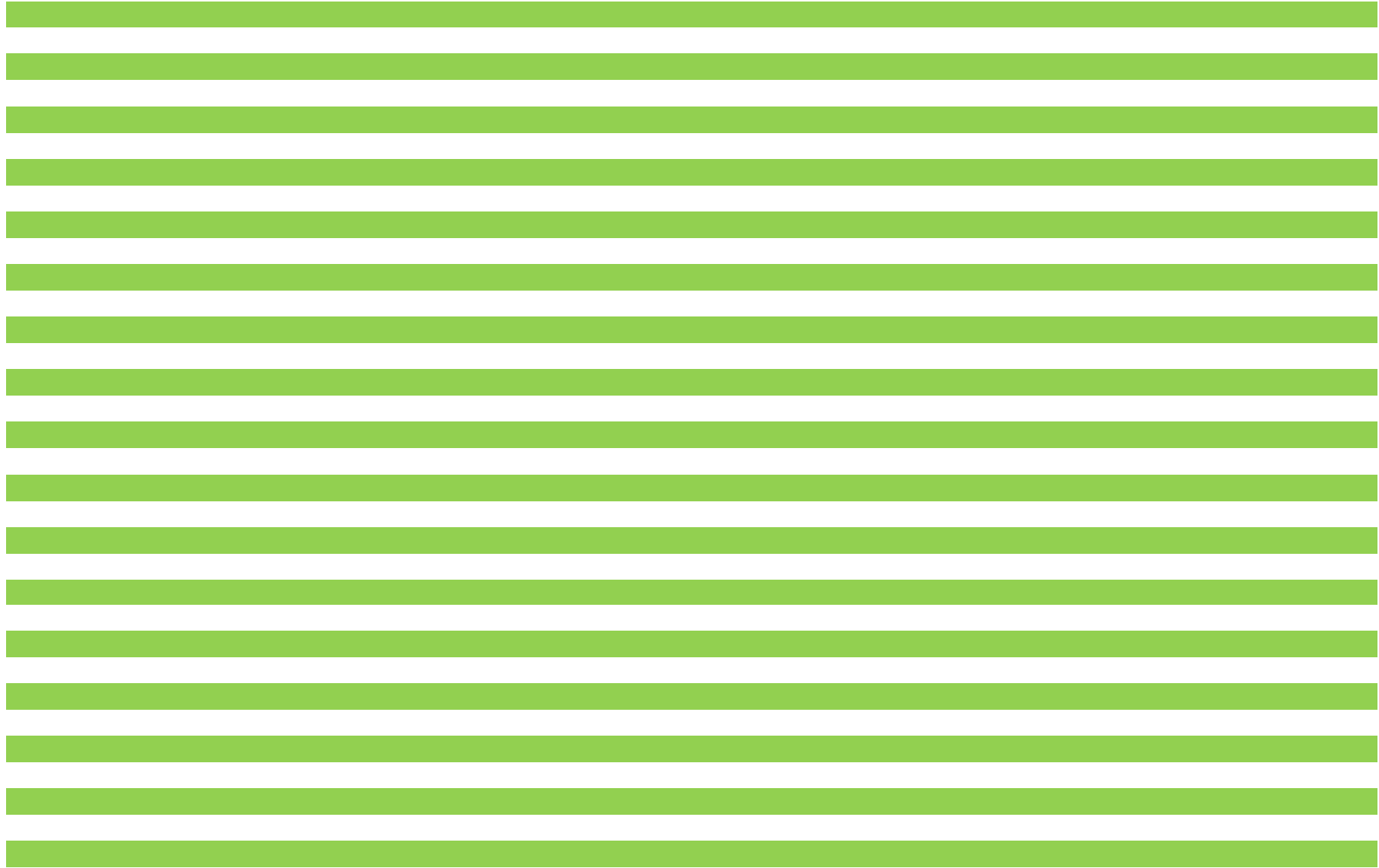
TAB]



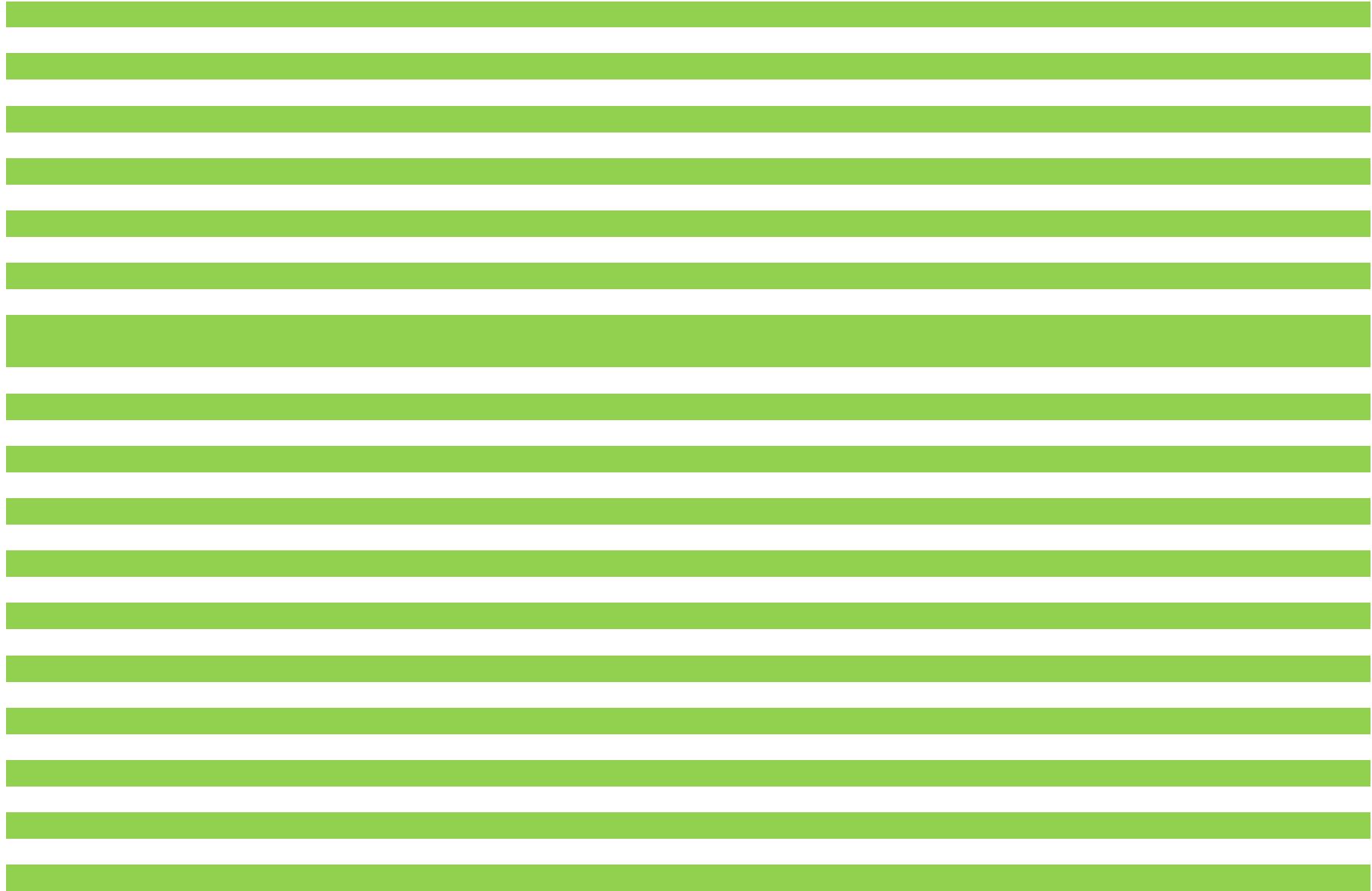
TAB]



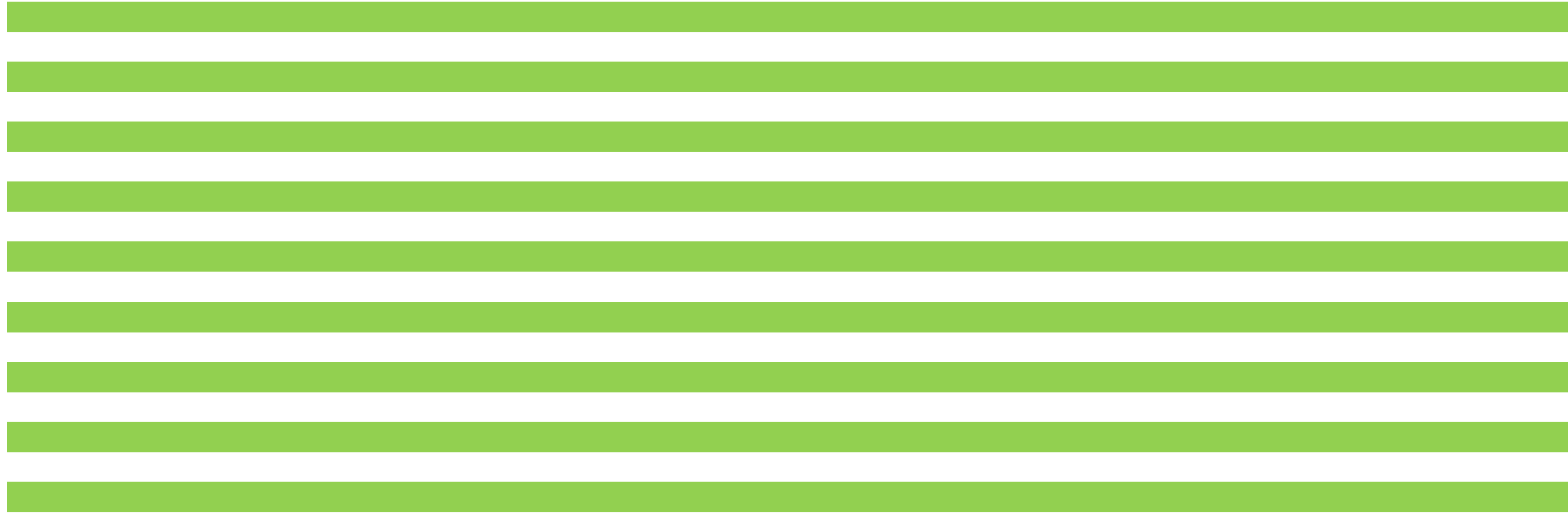
TAB]



TAB]



TAB]



TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

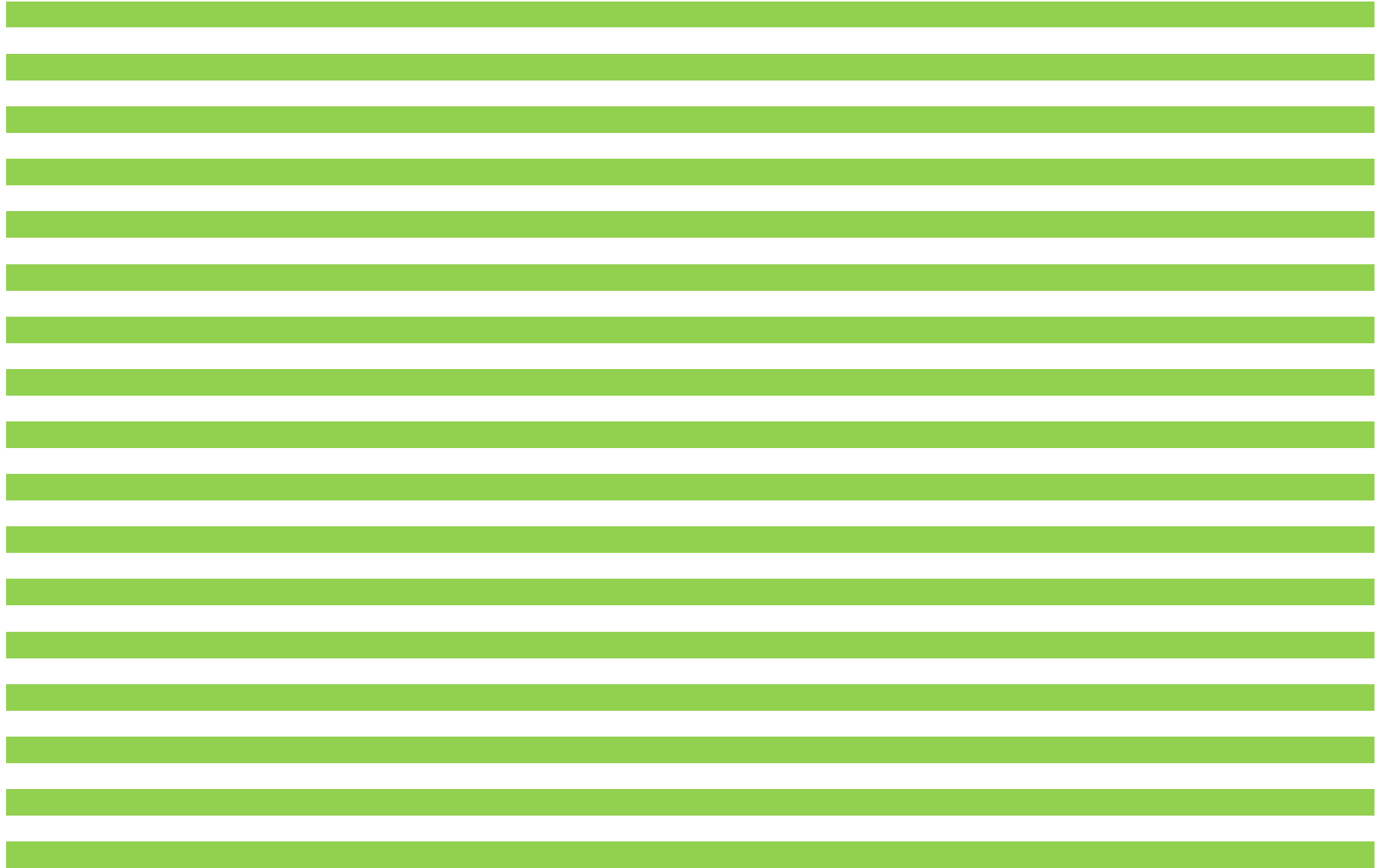
TAB]

TAB]

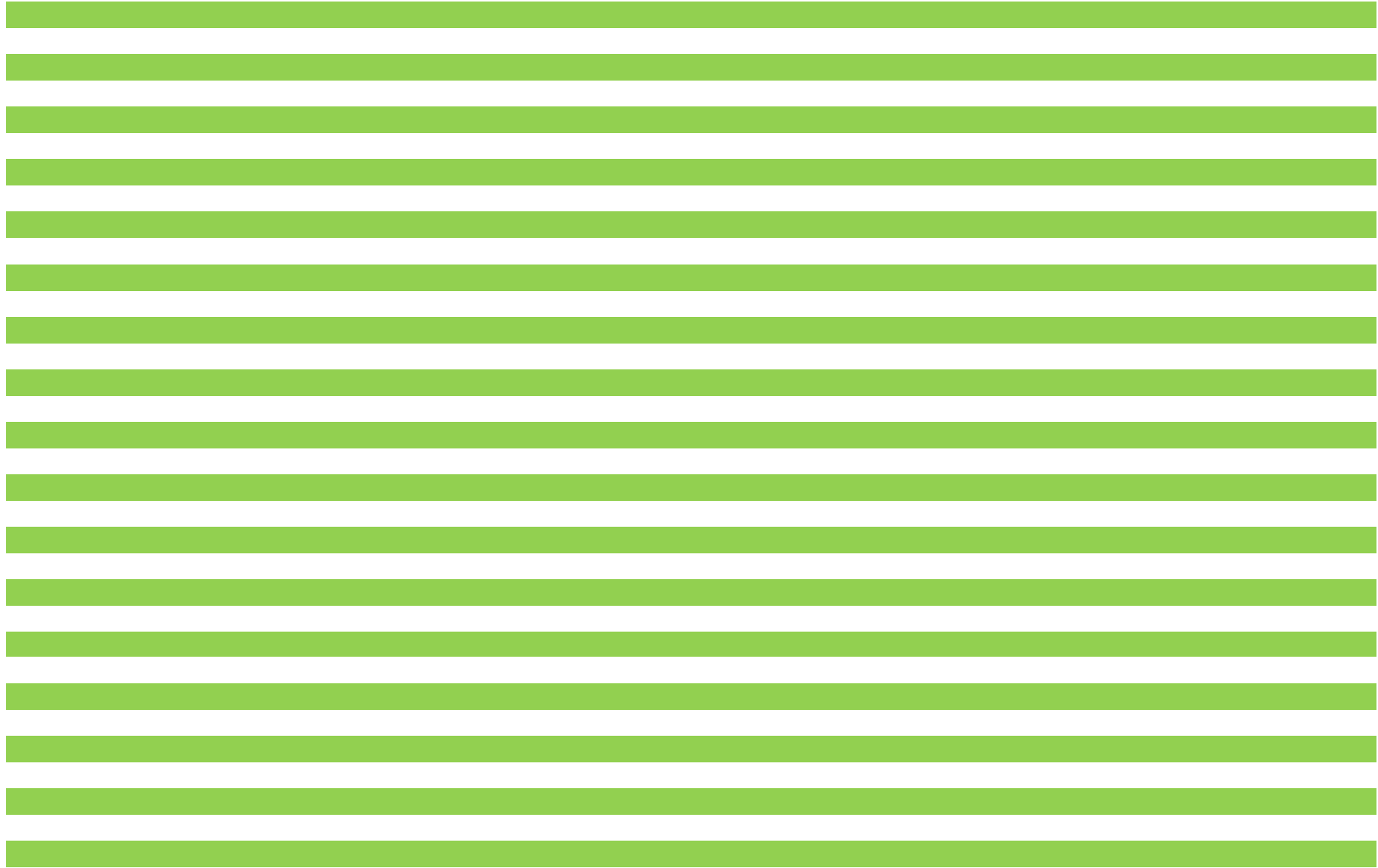
TAB]

TAB]

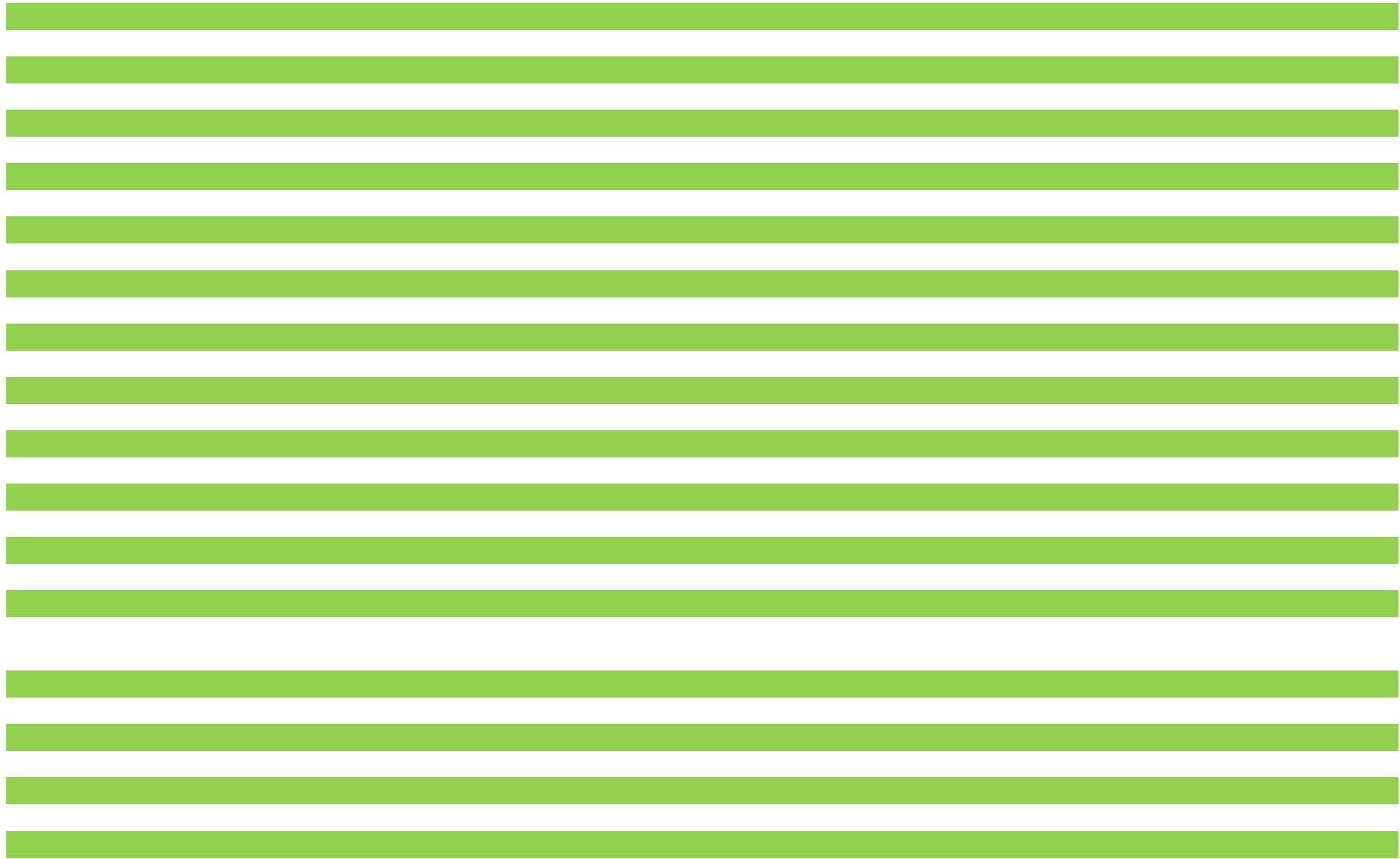
TAB]



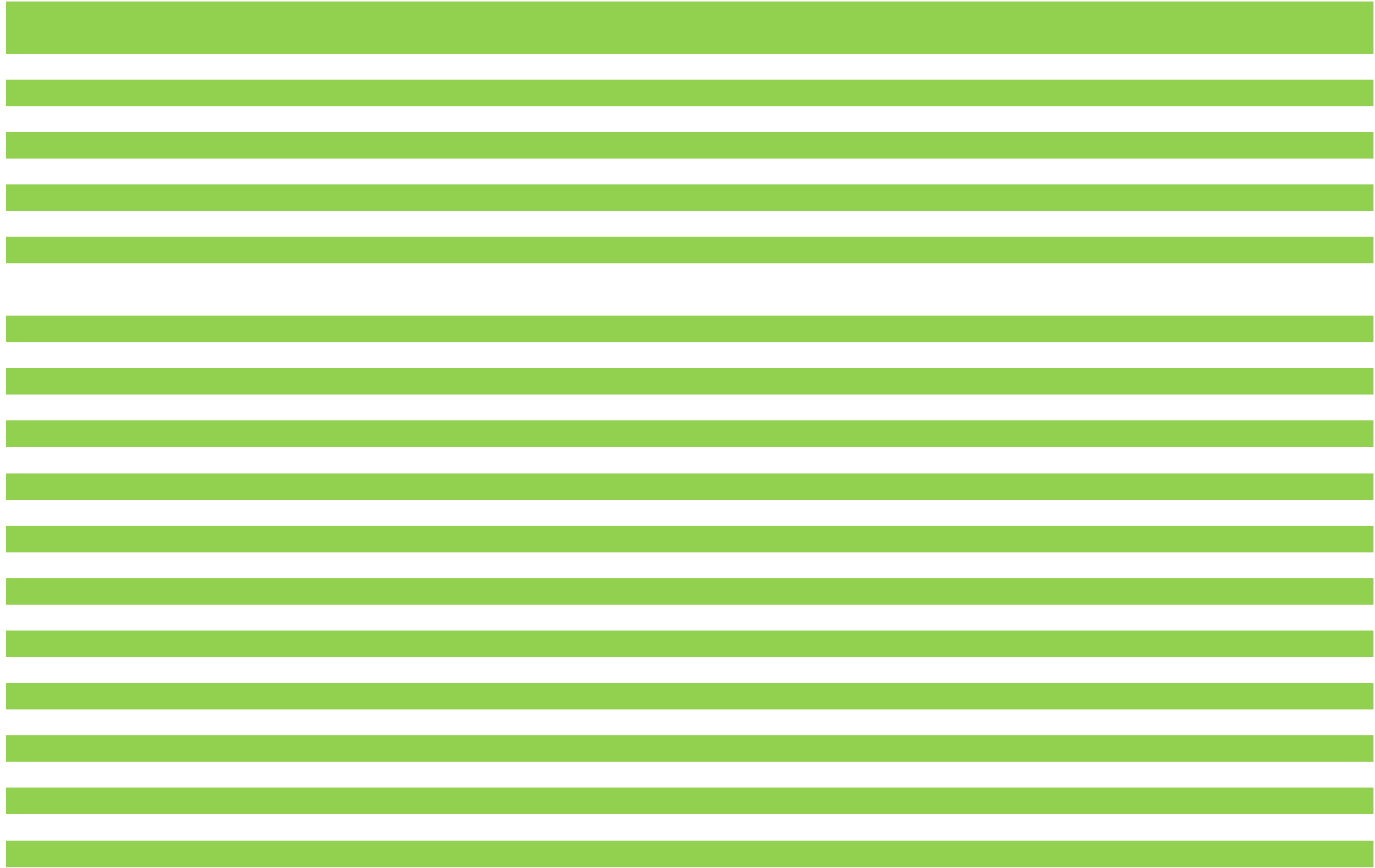
TAB]



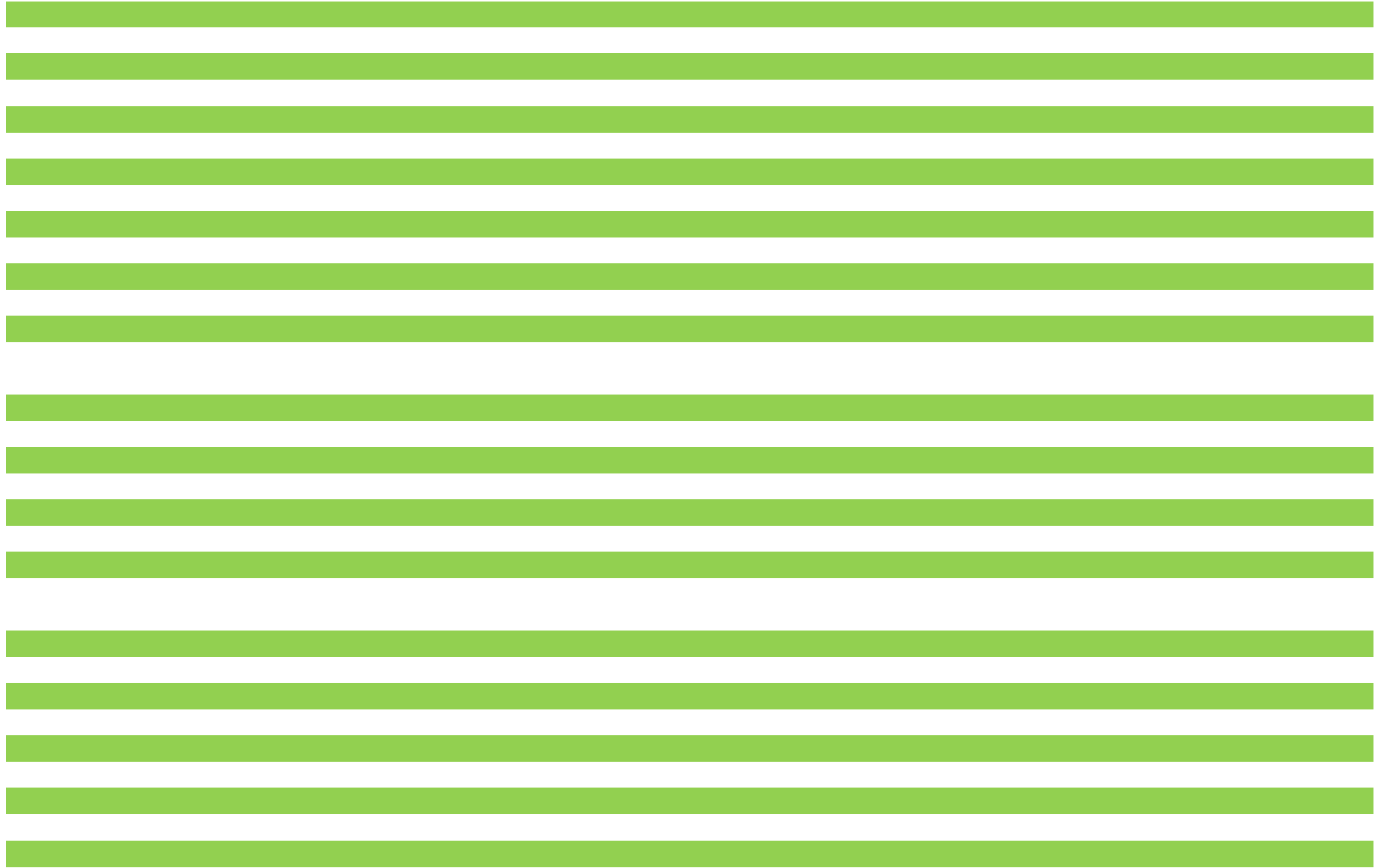
TAB]



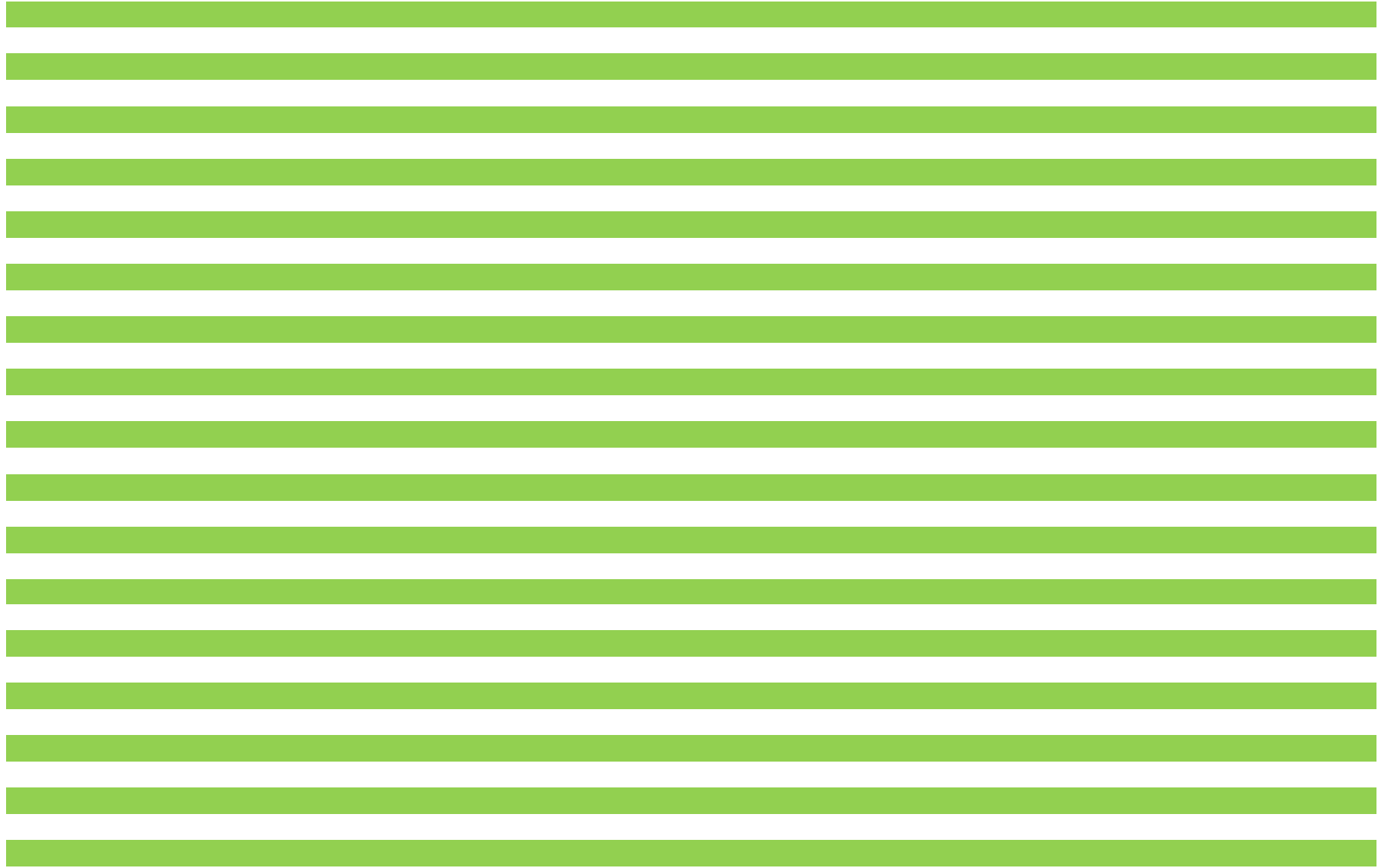
TAB]



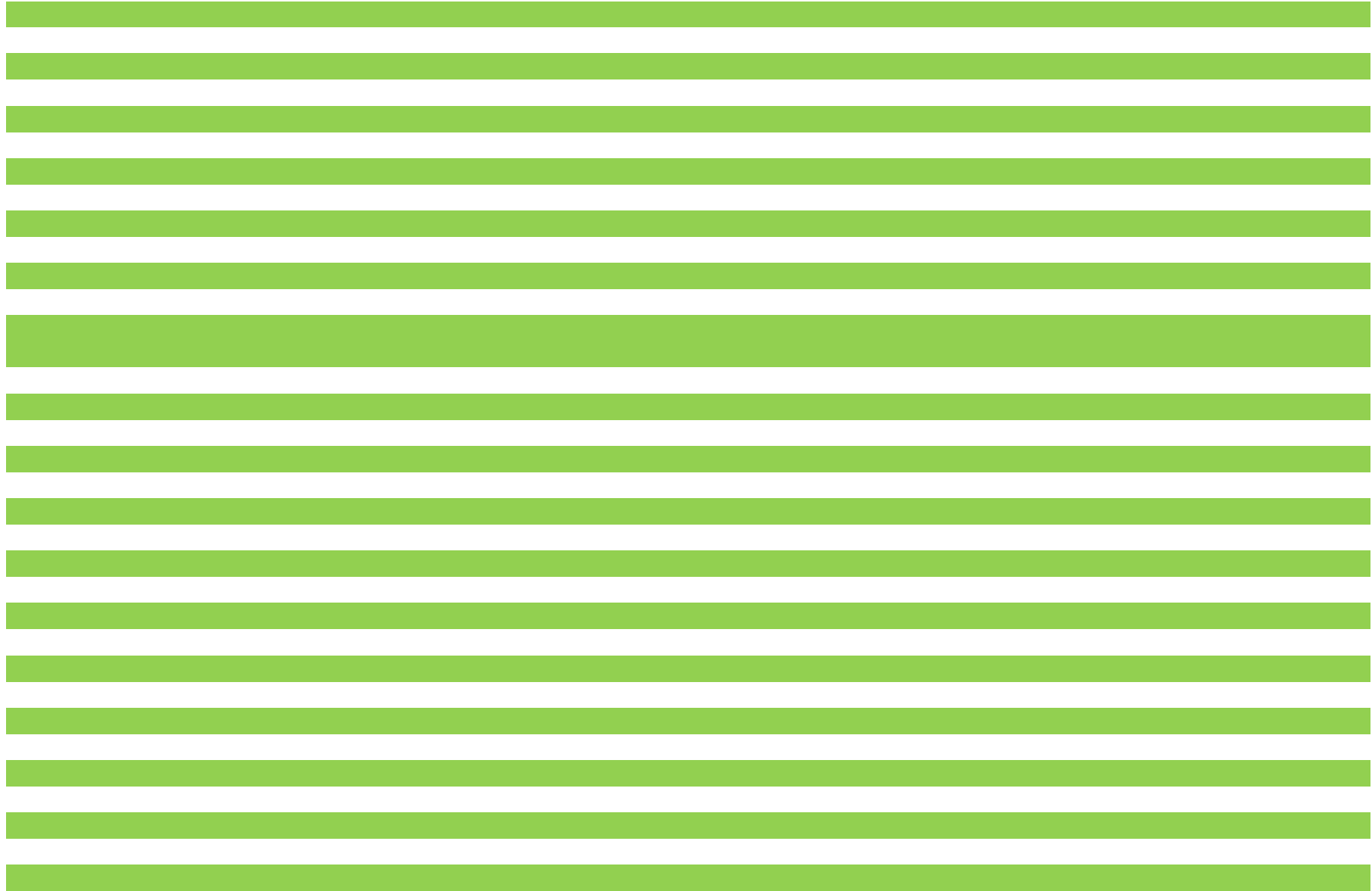
TAB]



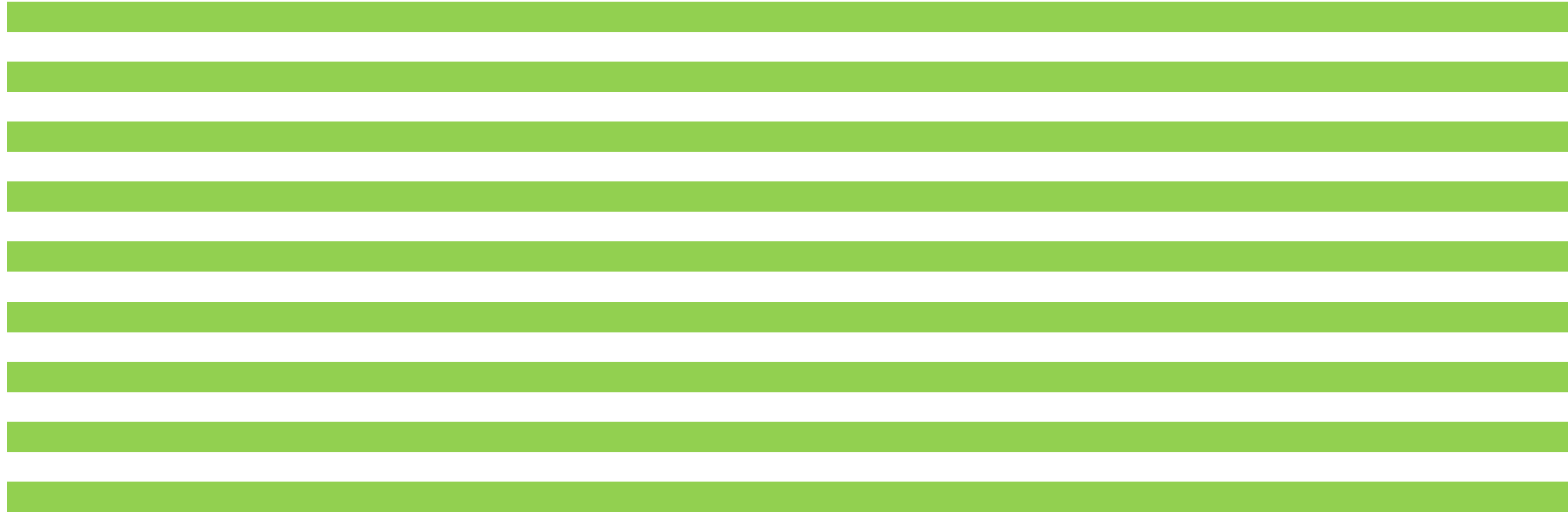
TAB]



TAB]



TAB]



TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

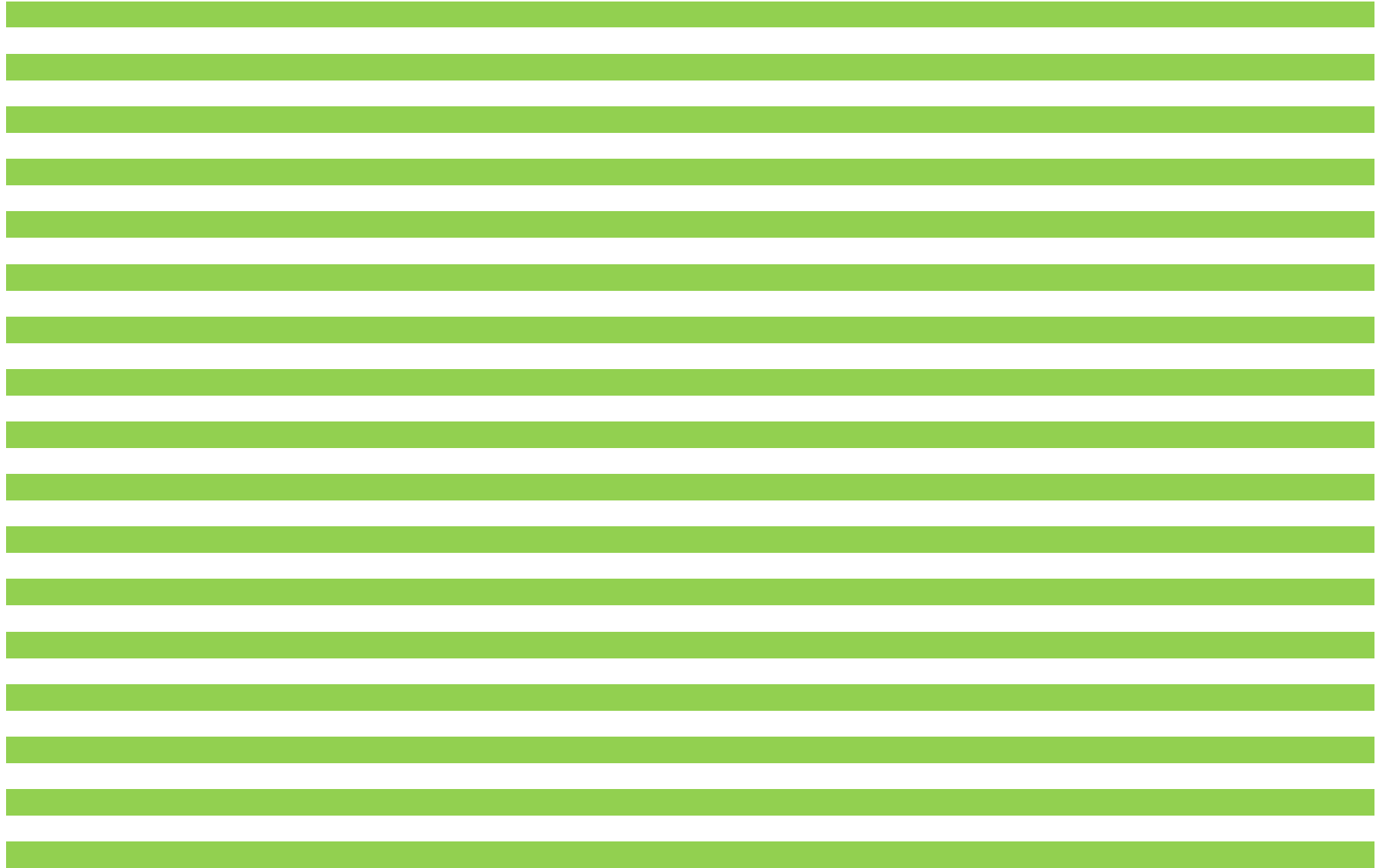
TAB]

TAB]

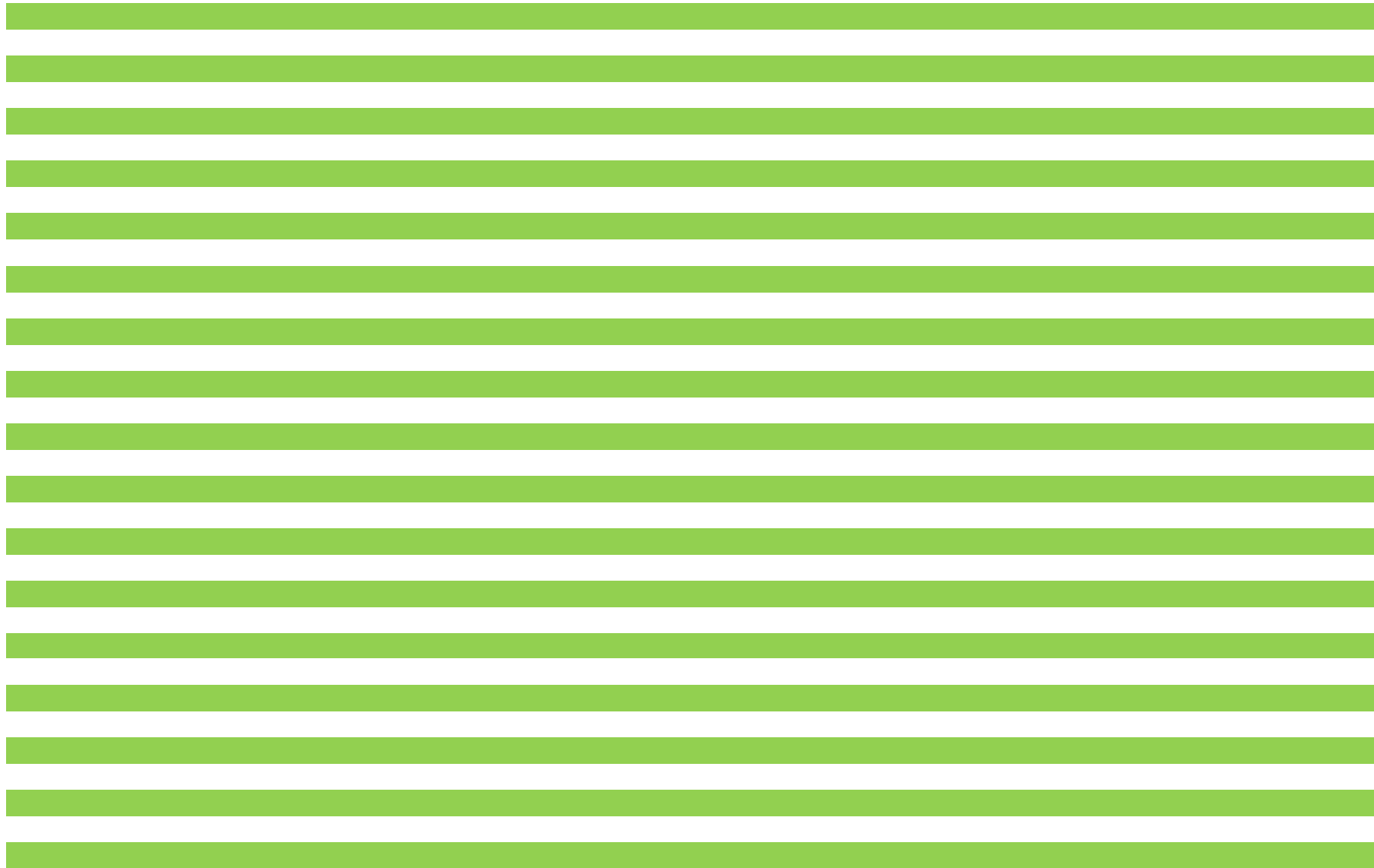
TAB]

TAB]

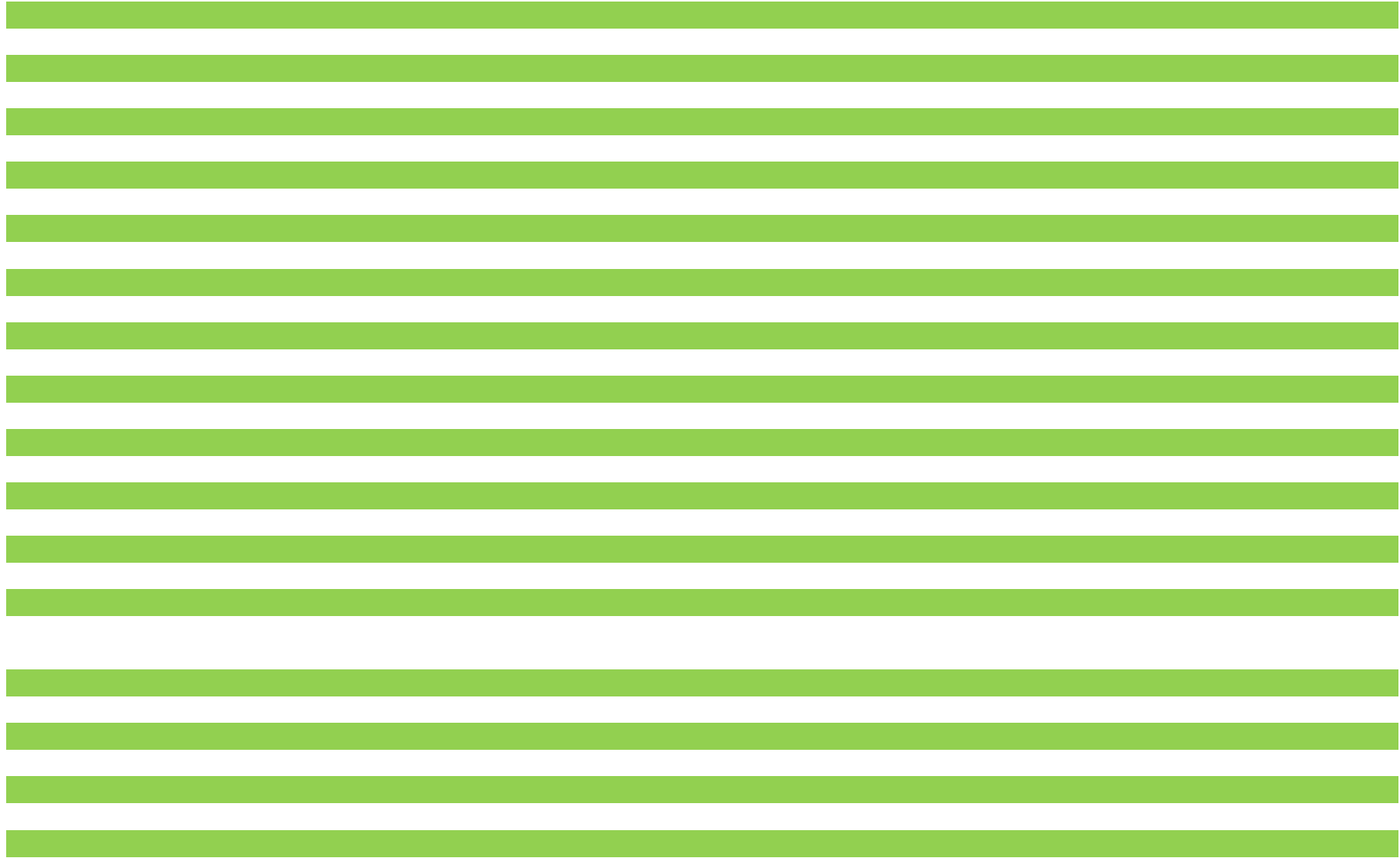
TAB]



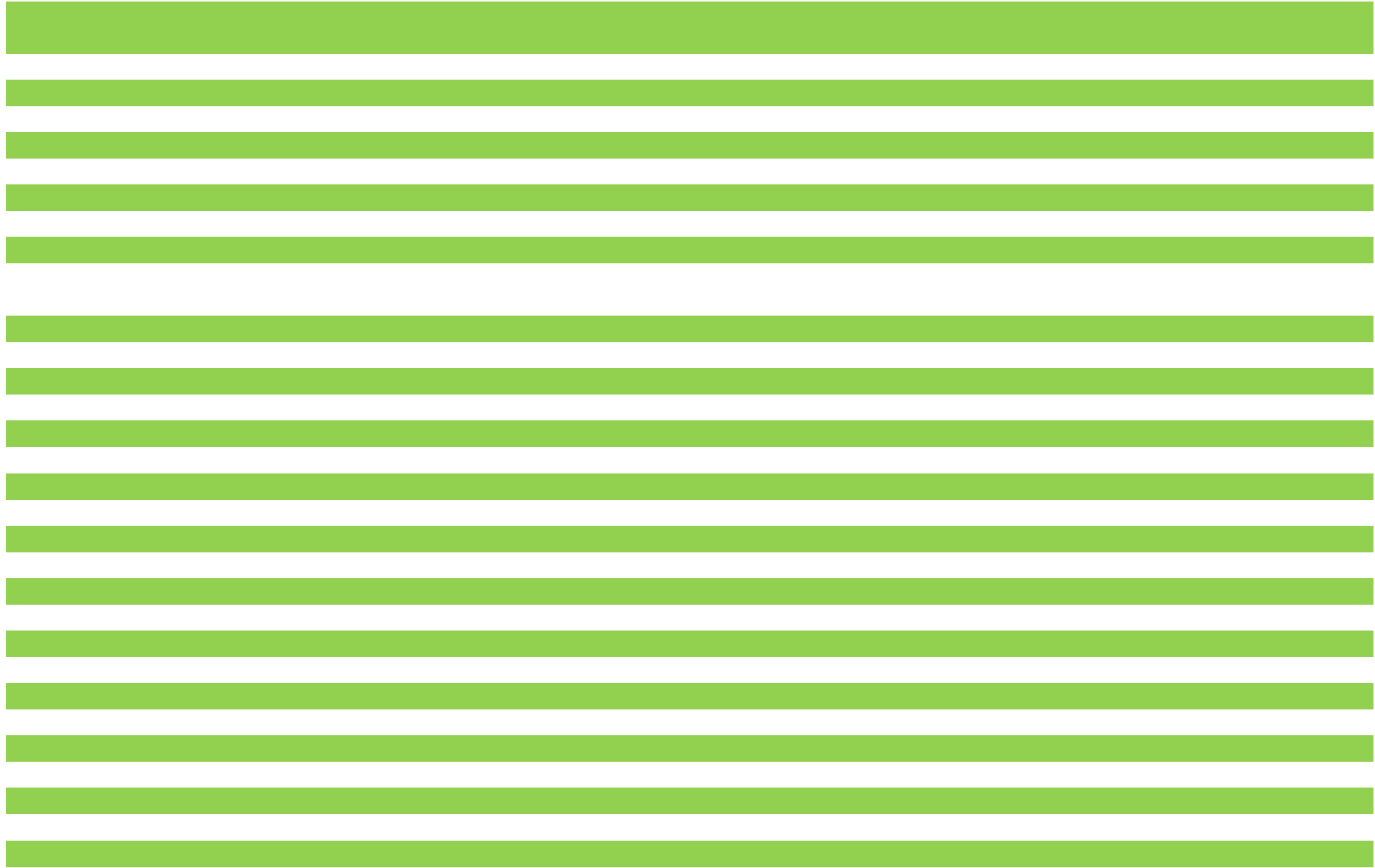
TAB]



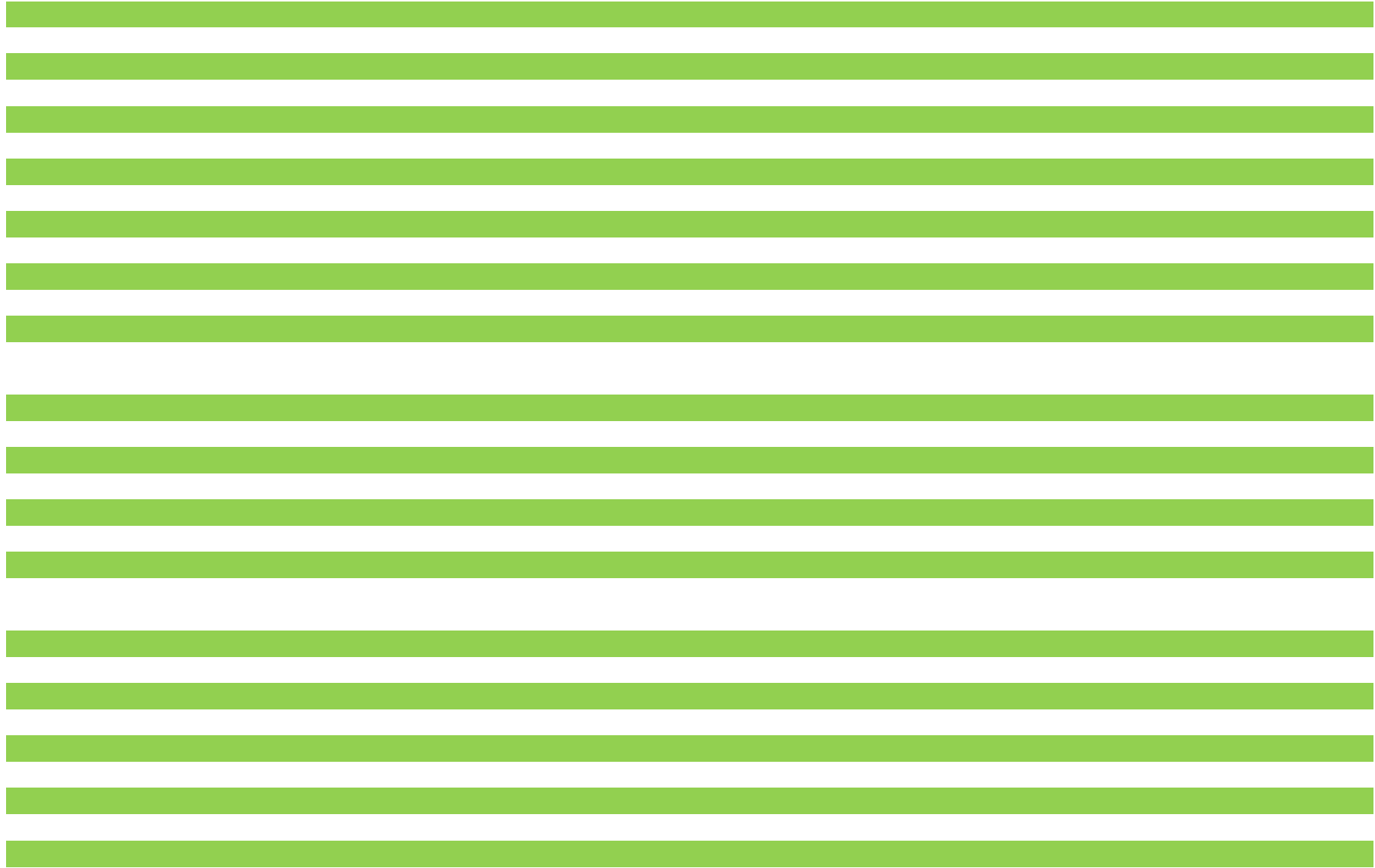
TAB]



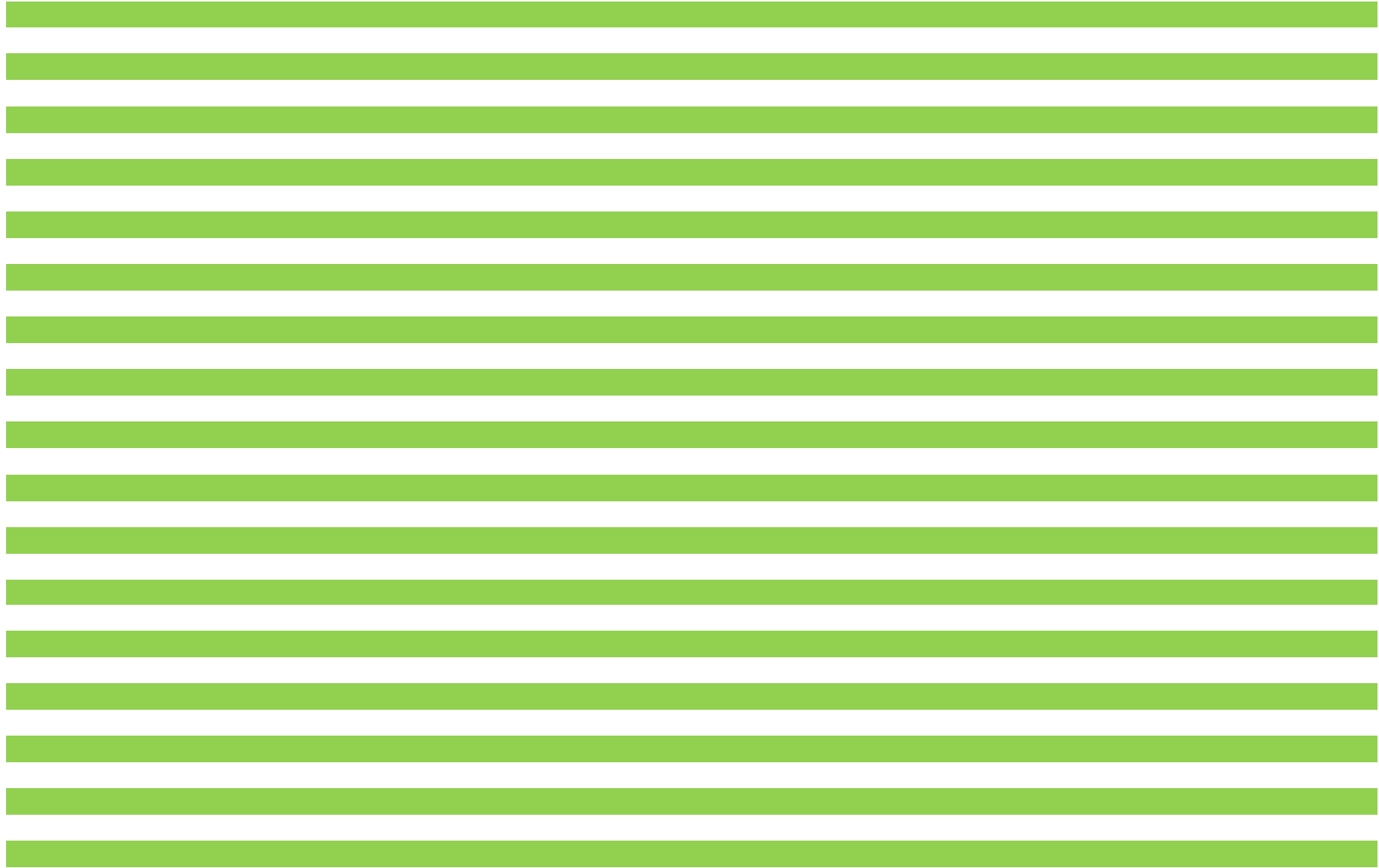
TAB]



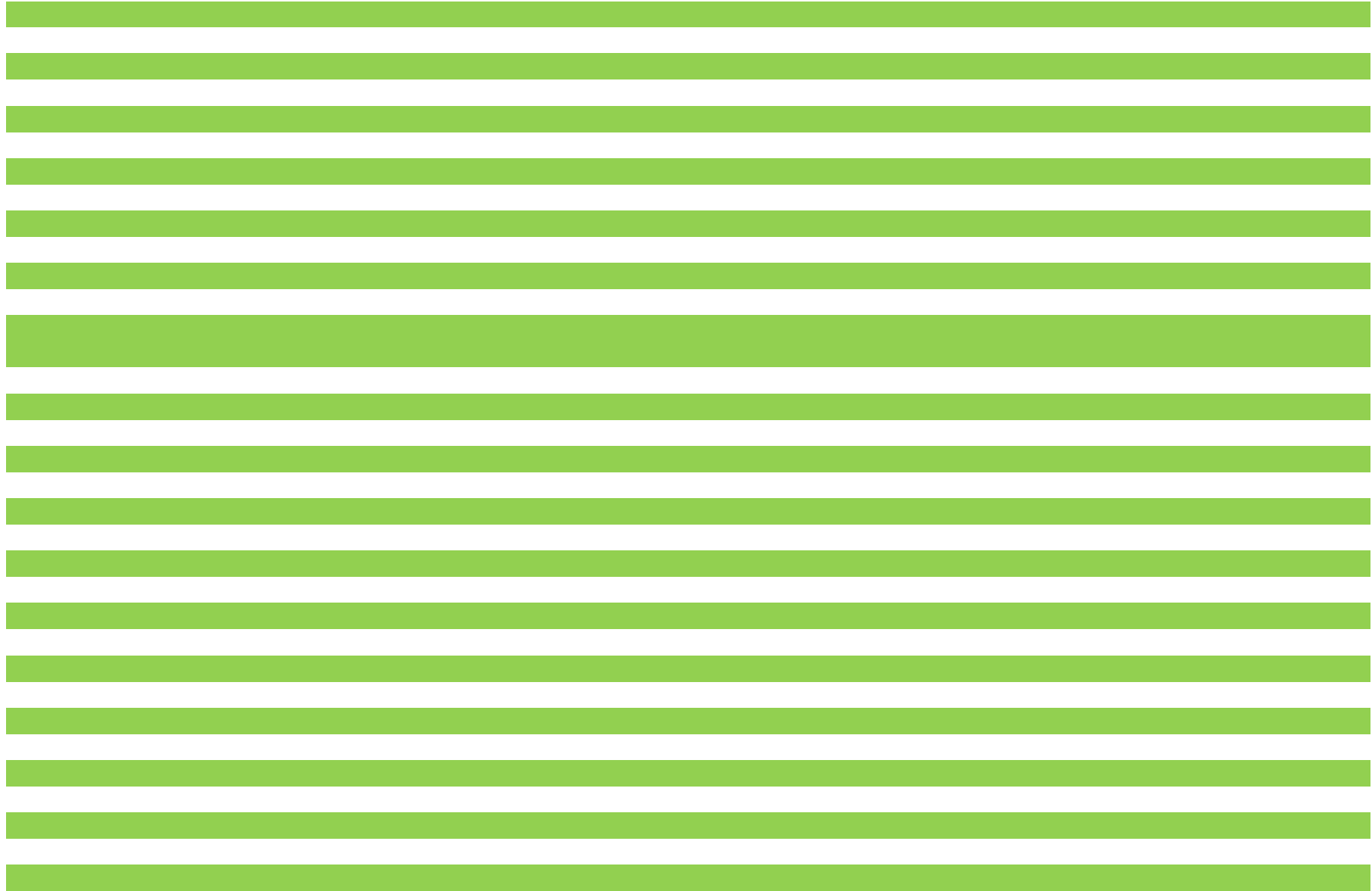
TAB]



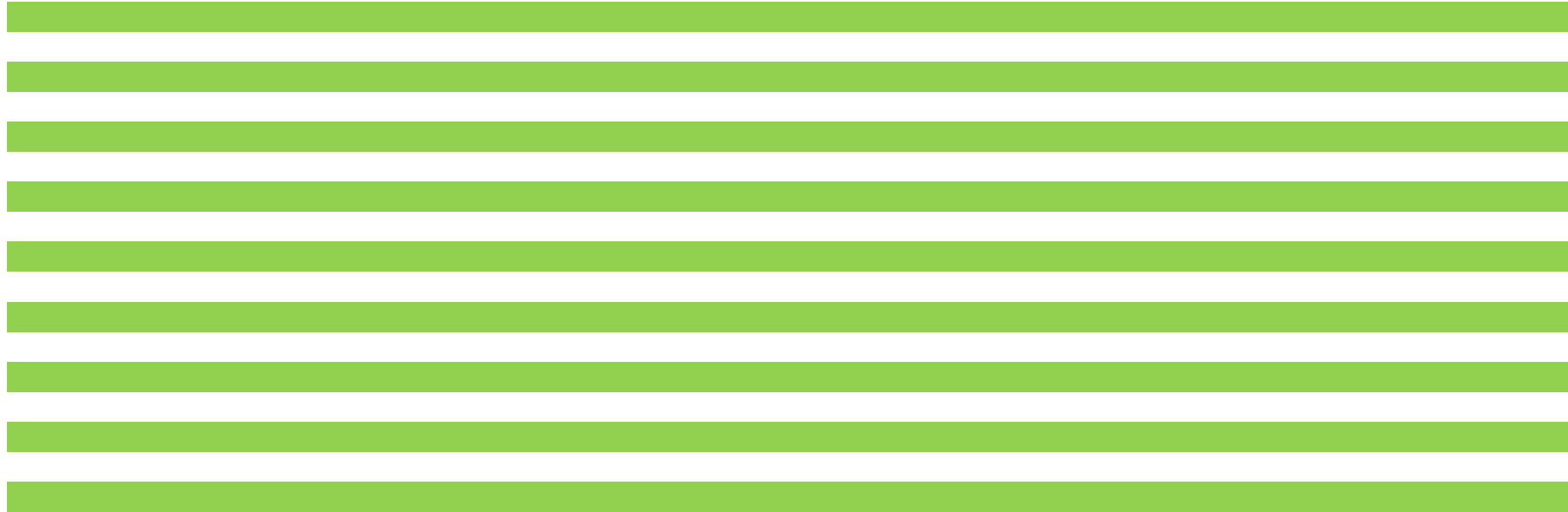
TAB]



TAB]



TAB]



TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

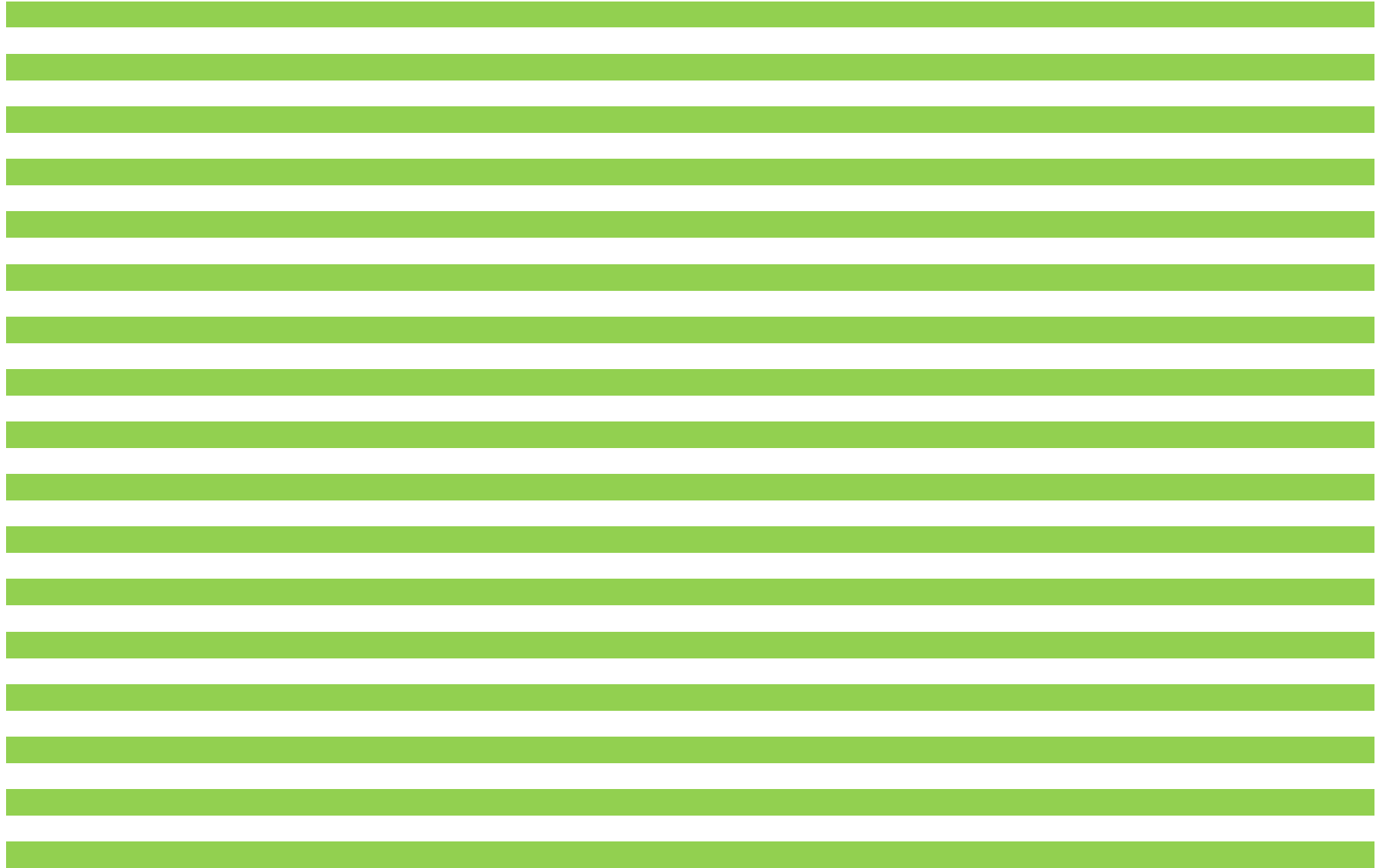
TAB]

TAB]

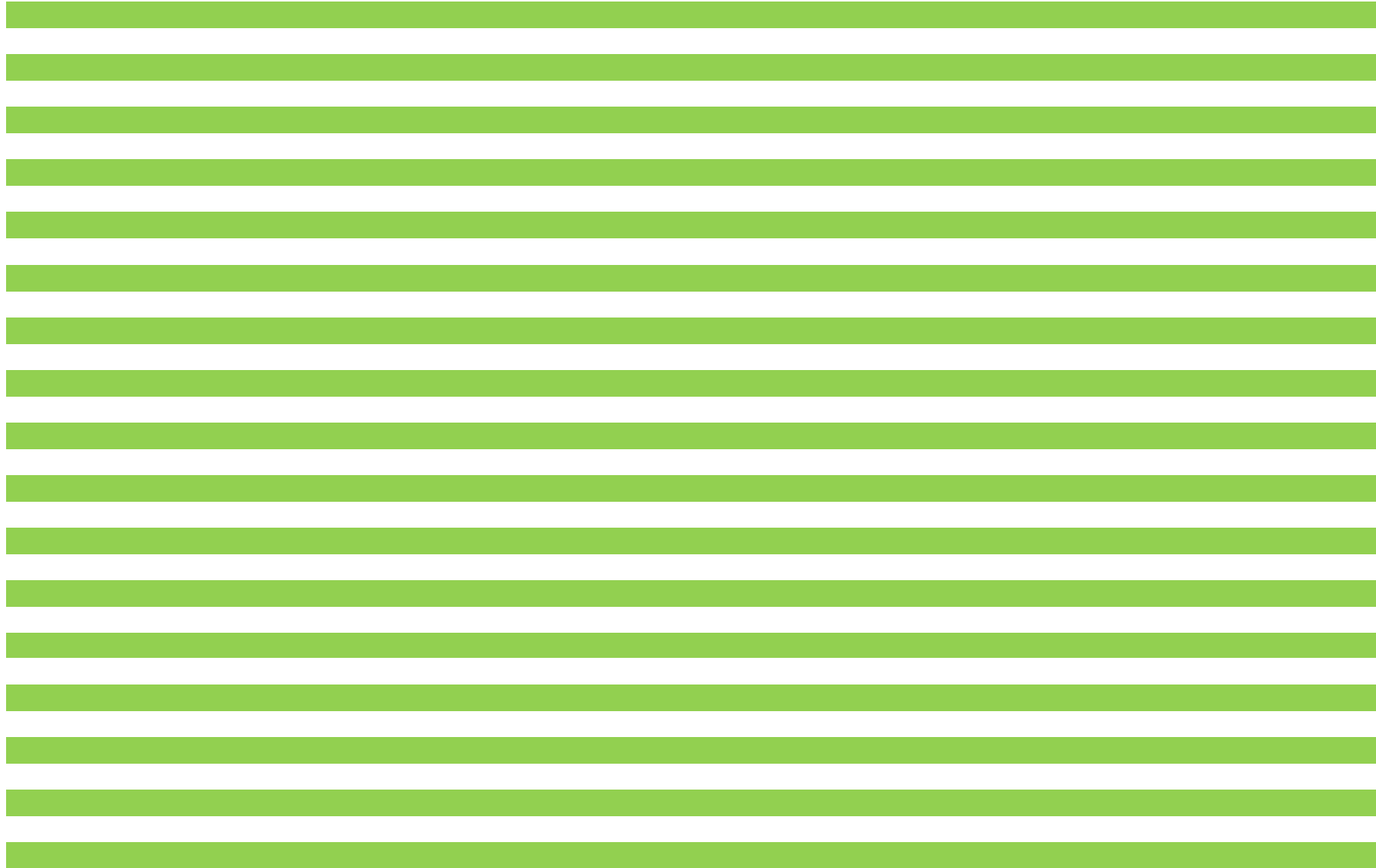
TAB]

TAB]

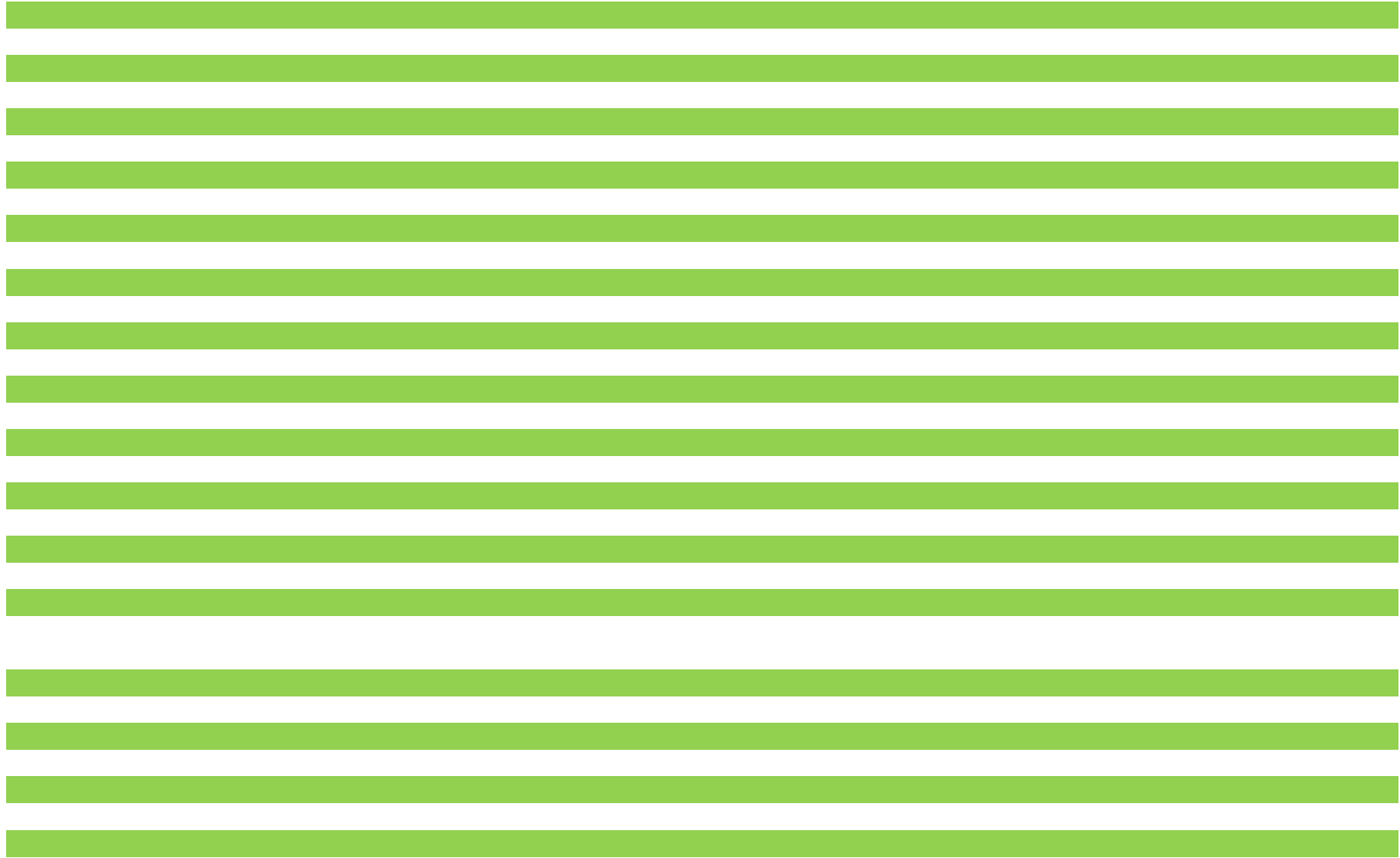
TAB]



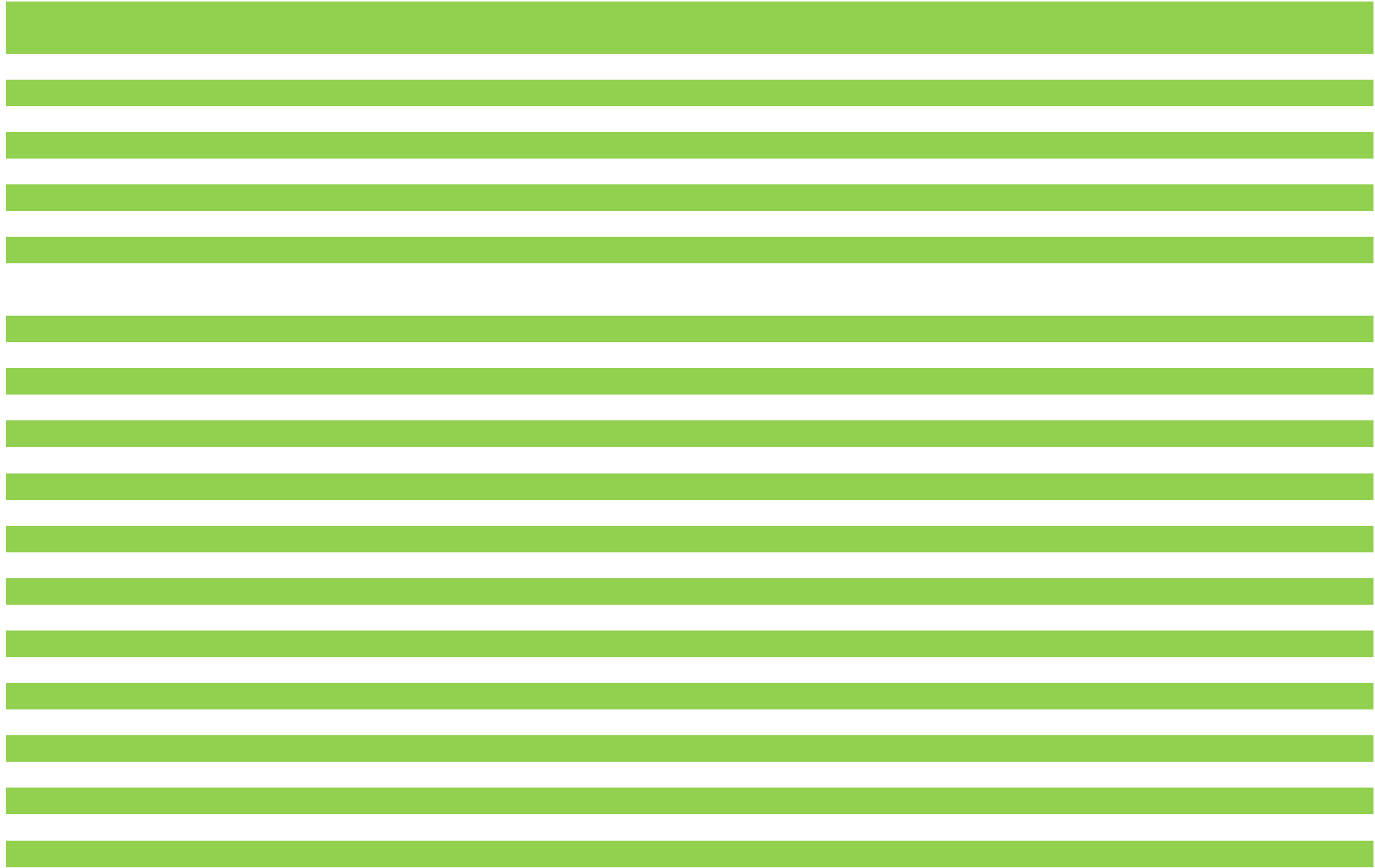
TAB]



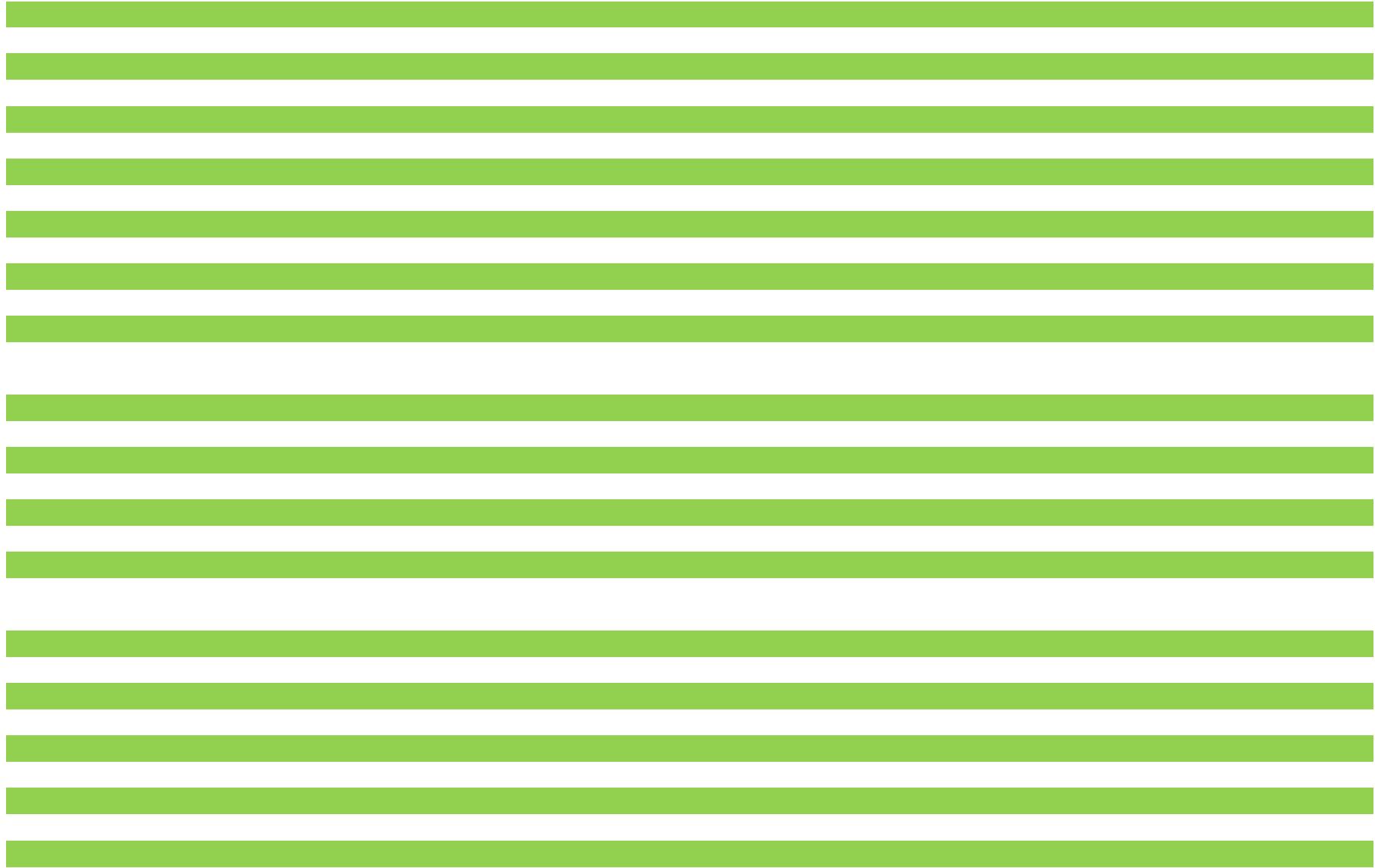
TAB]



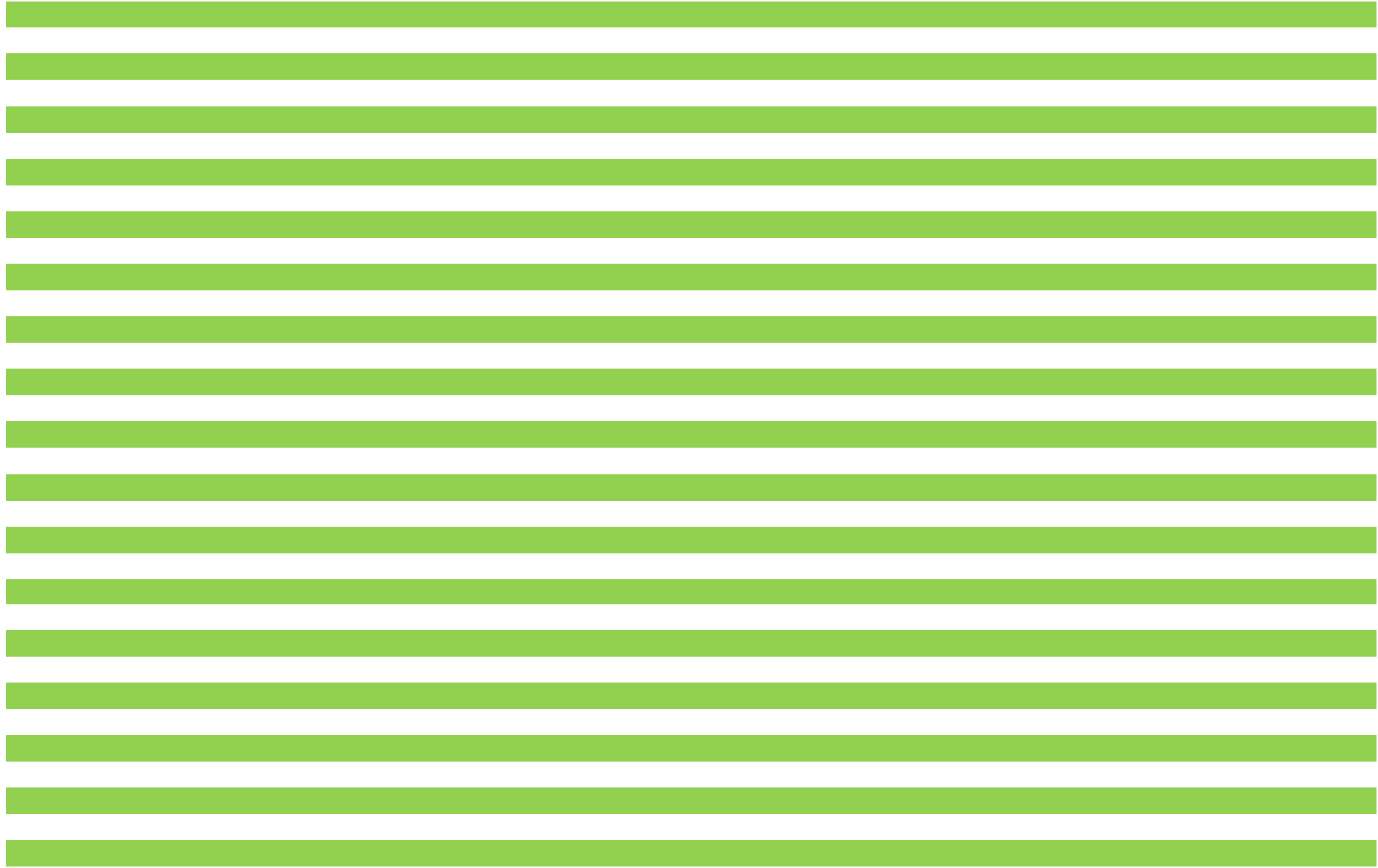
TAB]



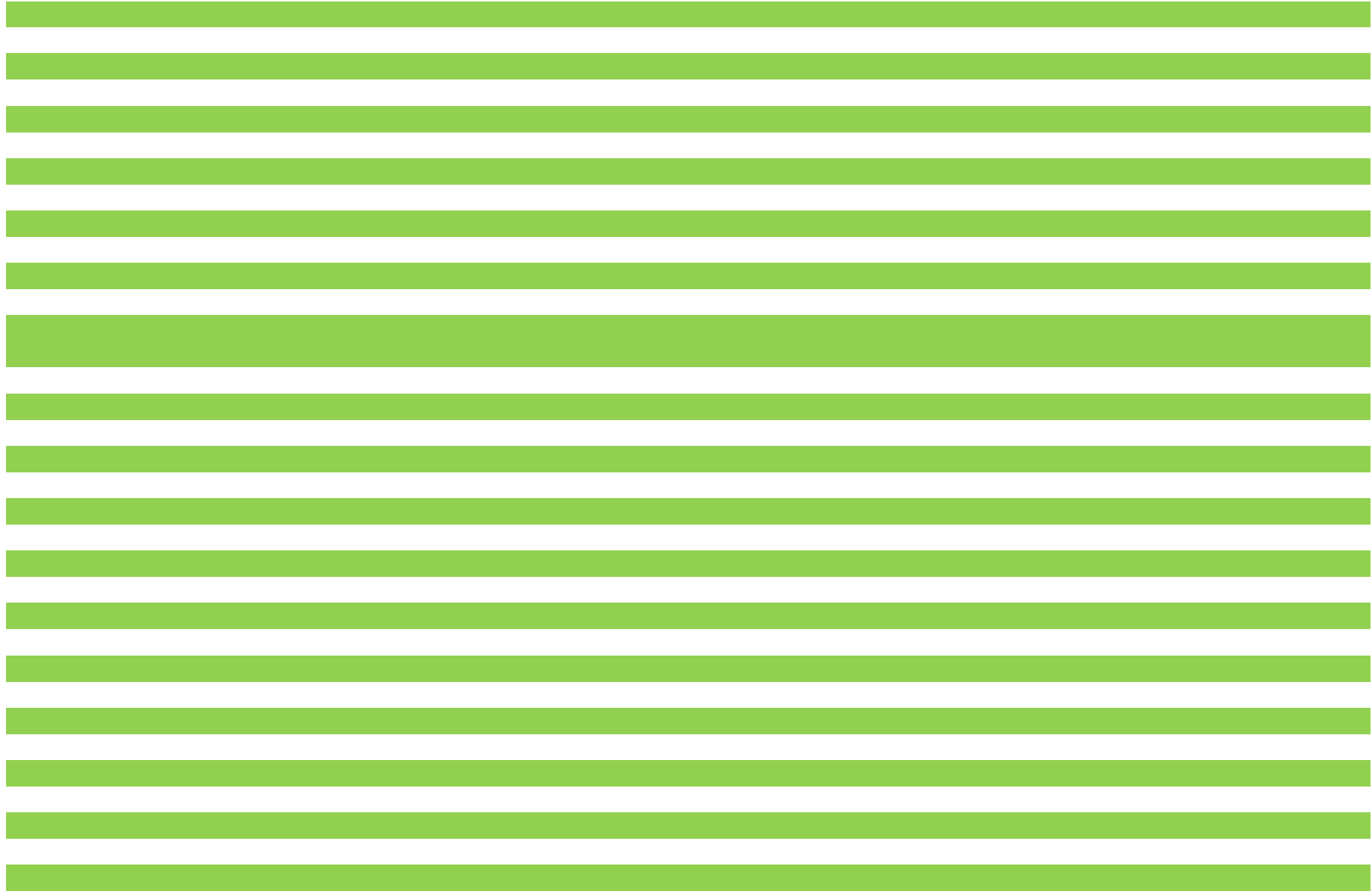
TAB]



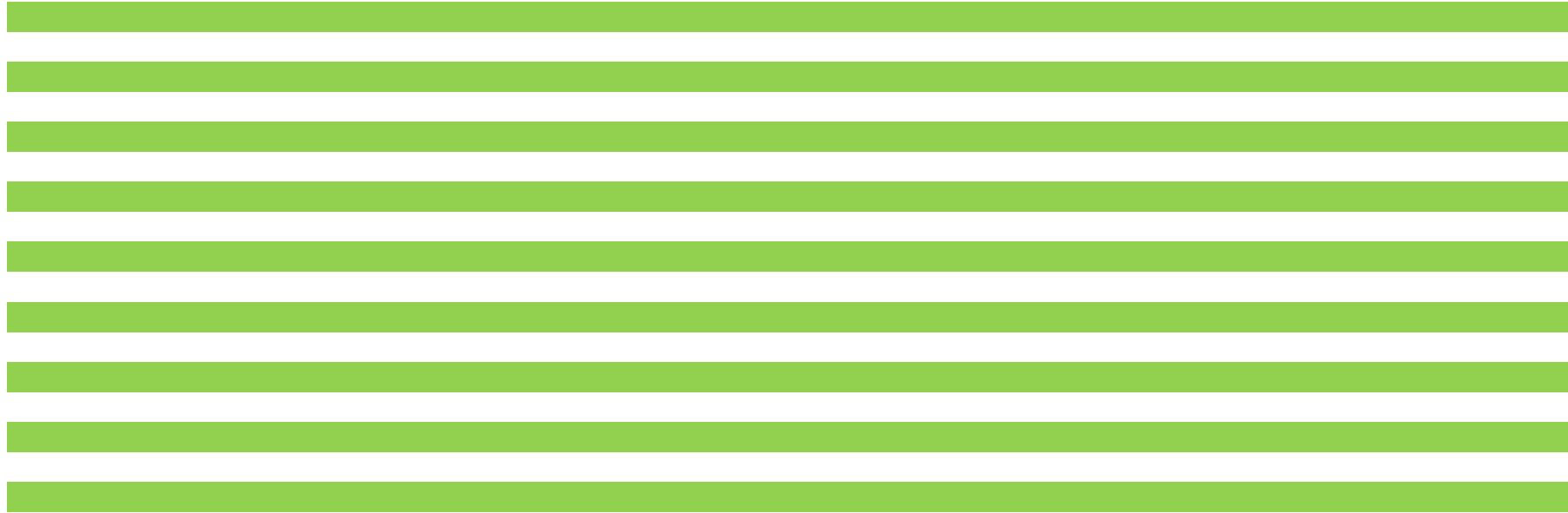
TAB]



TAB]



TAB]



TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

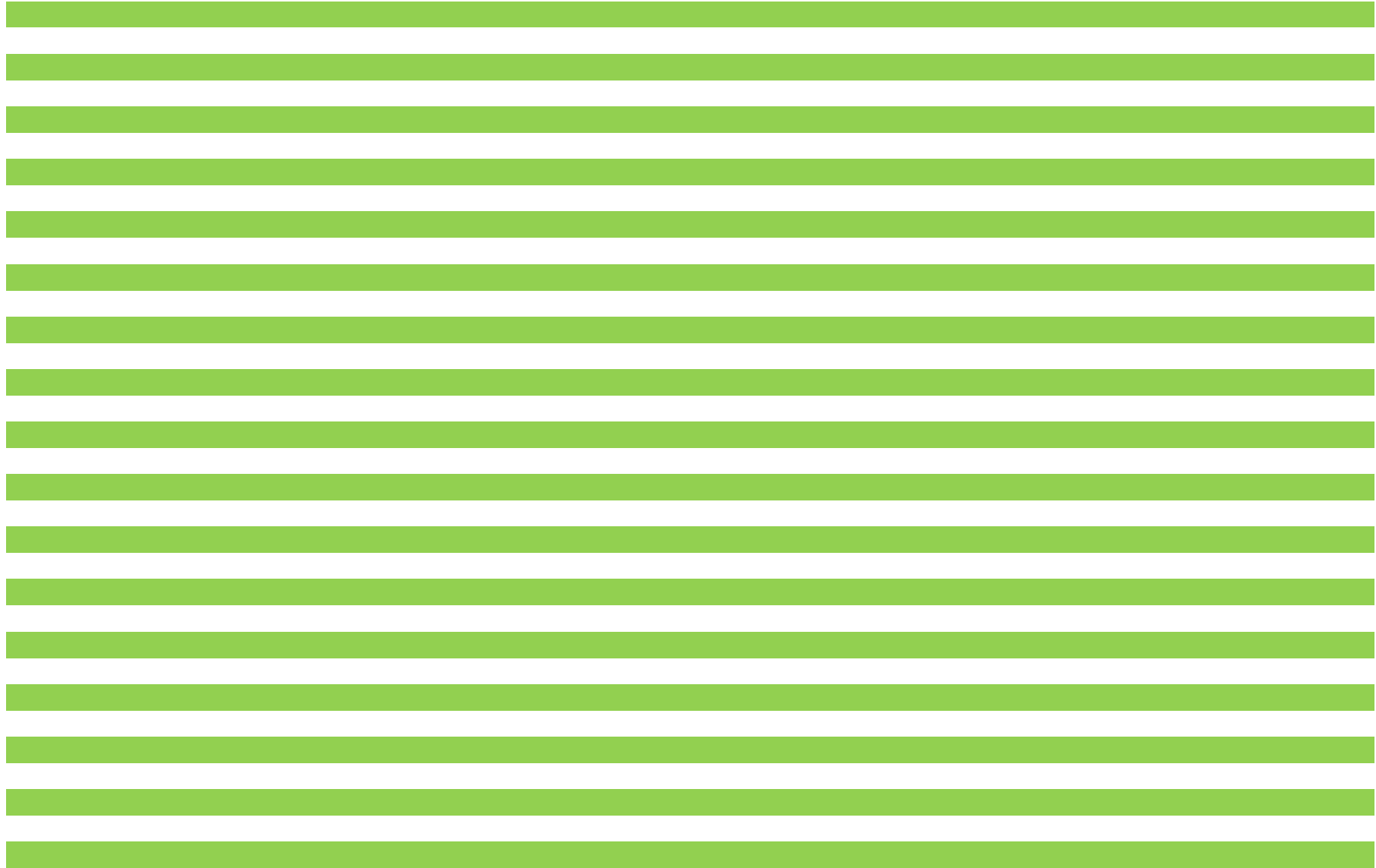
TAB]

TAB]

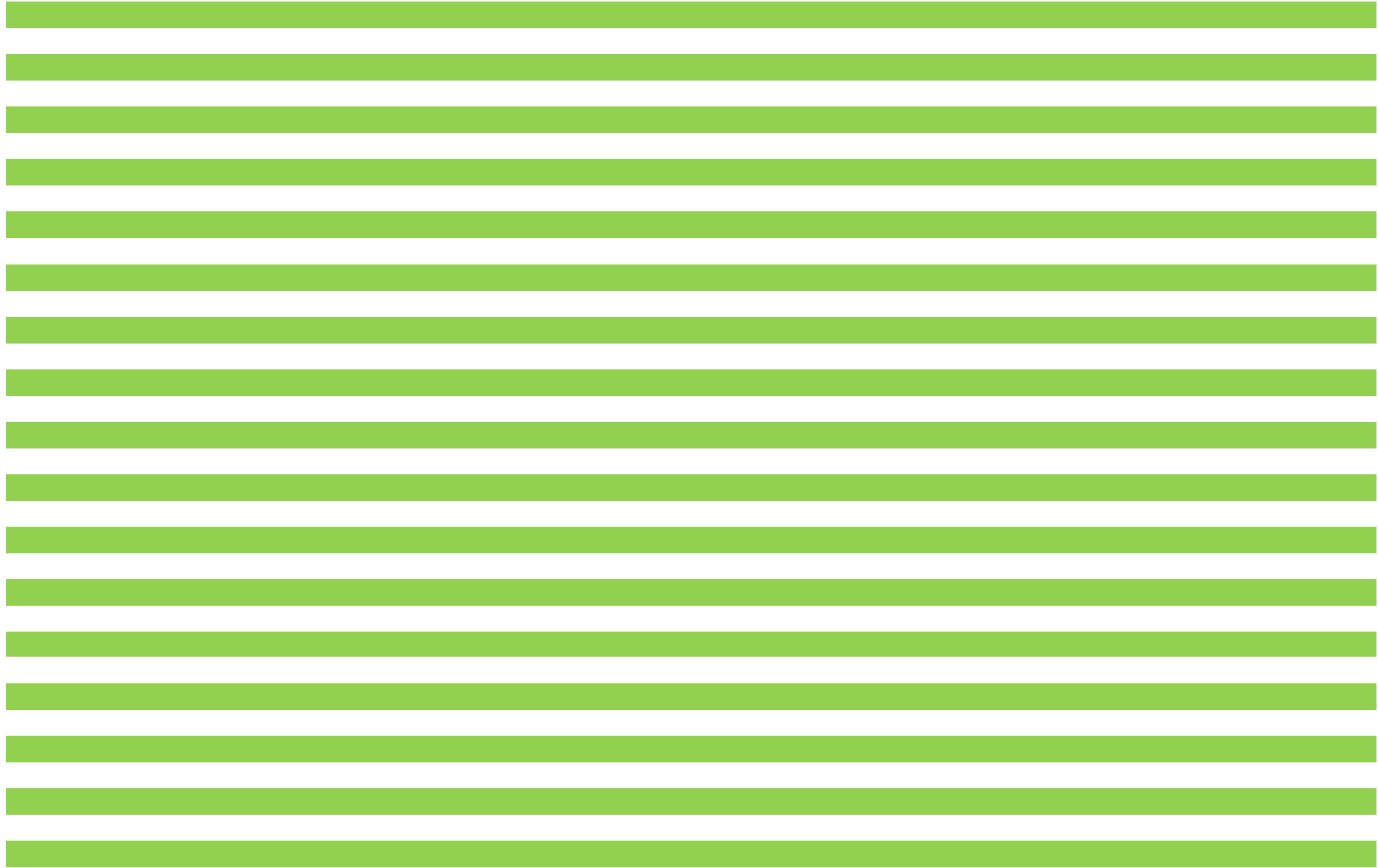
TAB]

TAB]

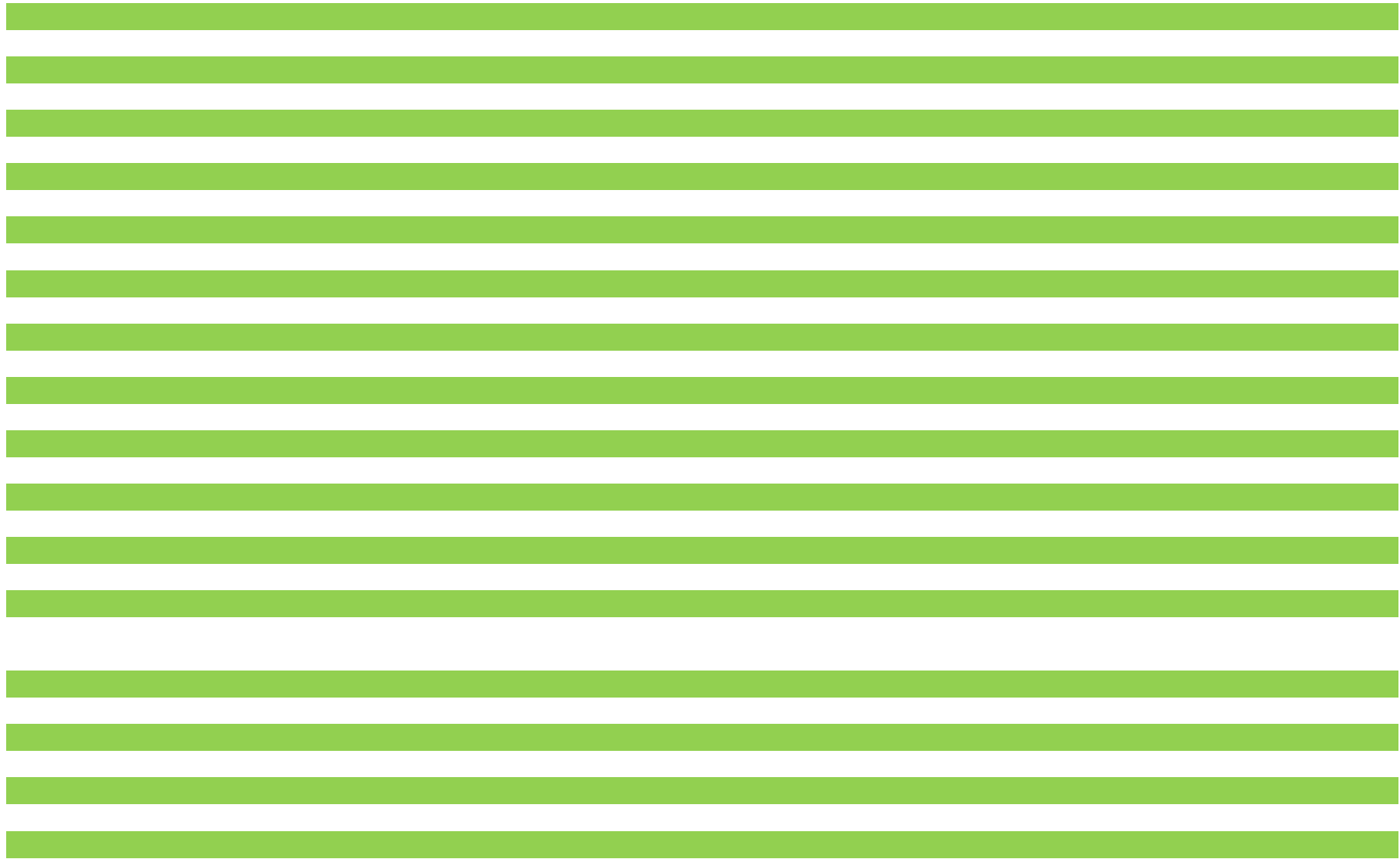
TAB]



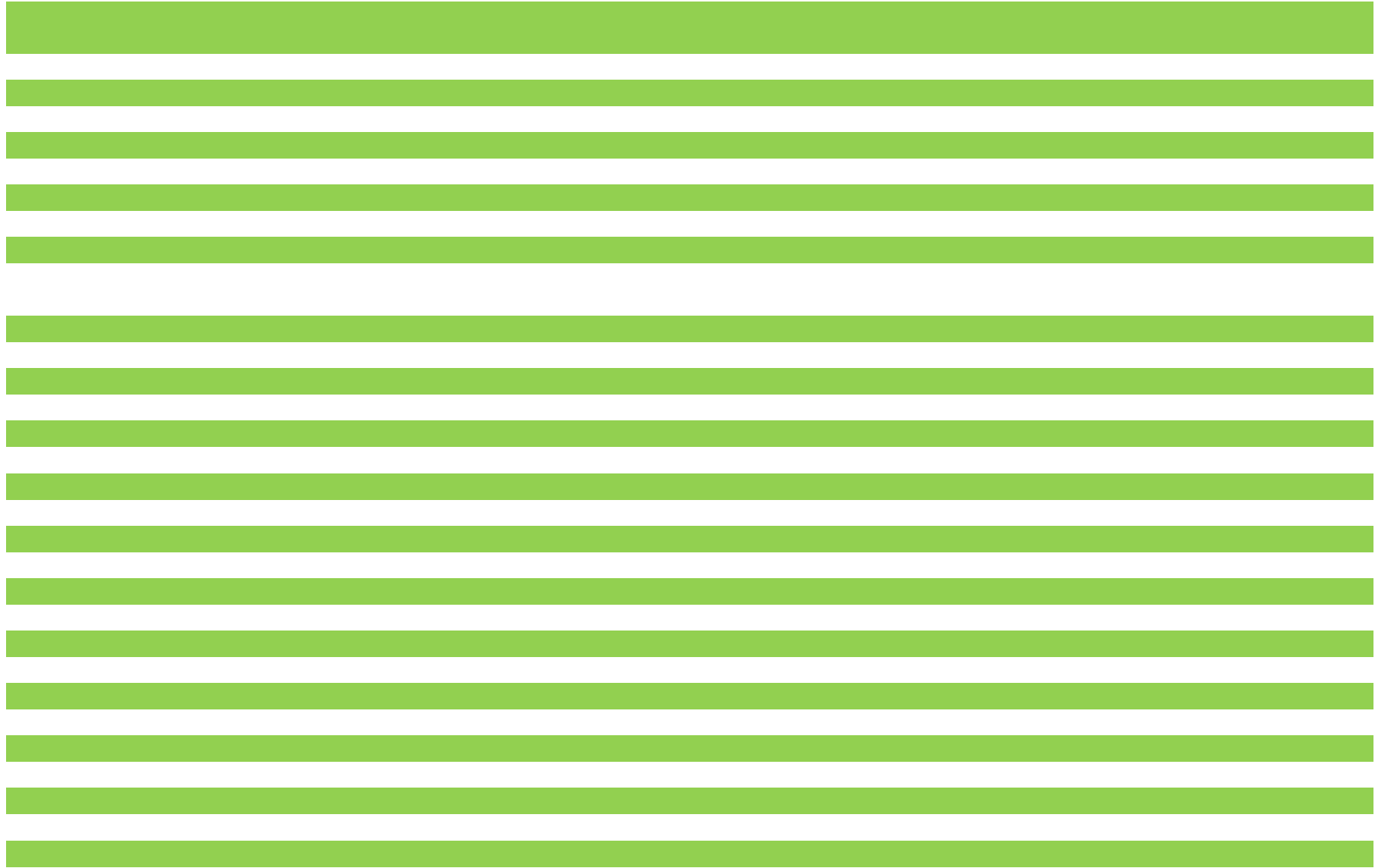
TAB]



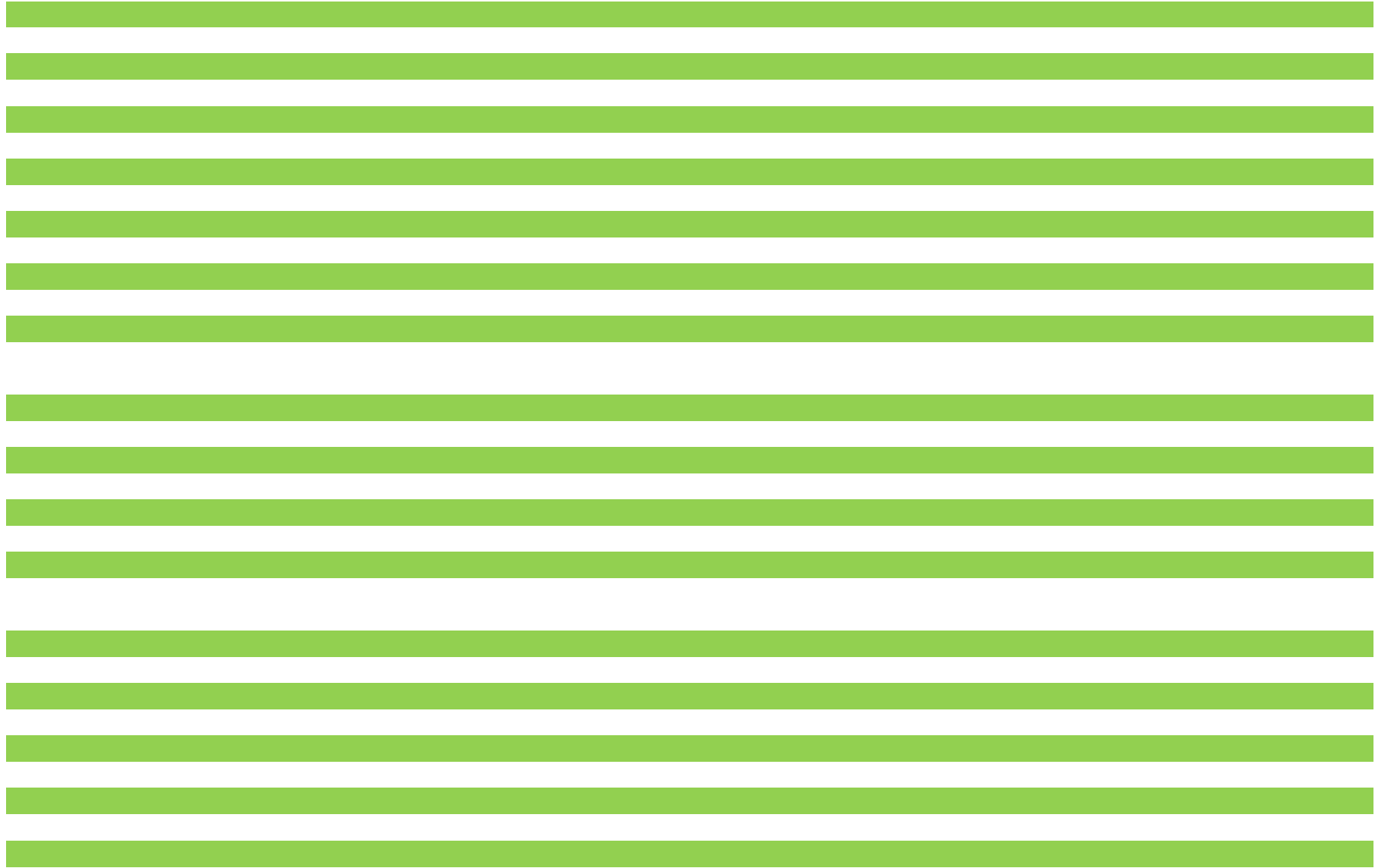
TAB]



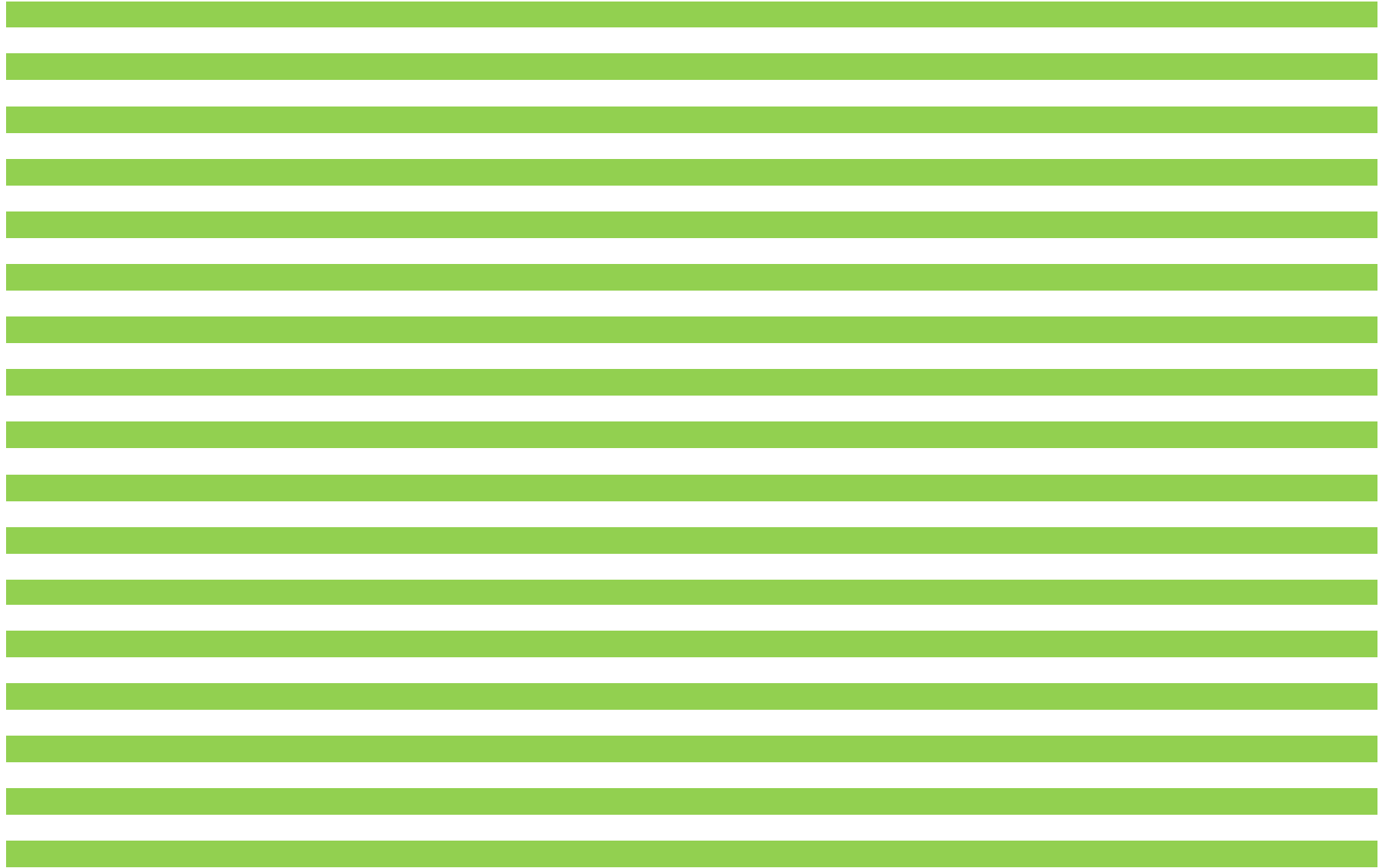
TAB]



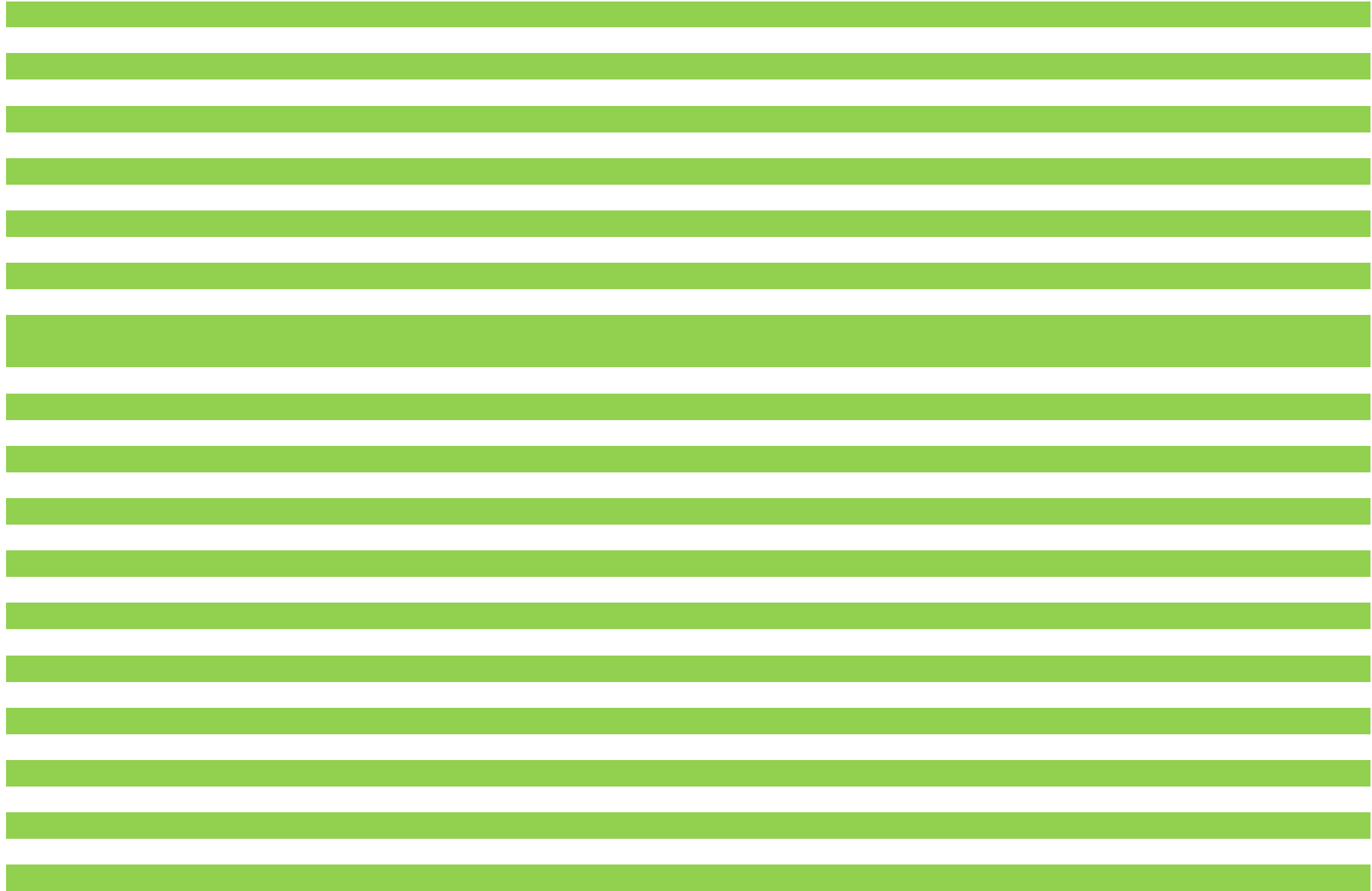
TAB]



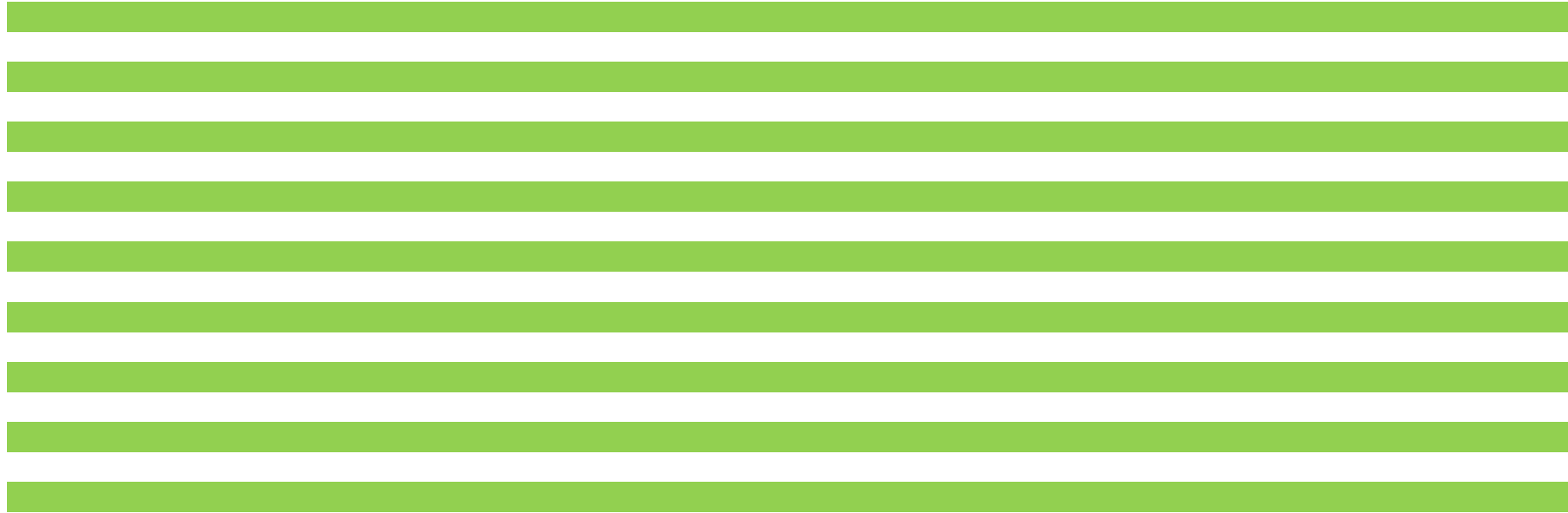
TAB]



TAB]



TAB]



TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

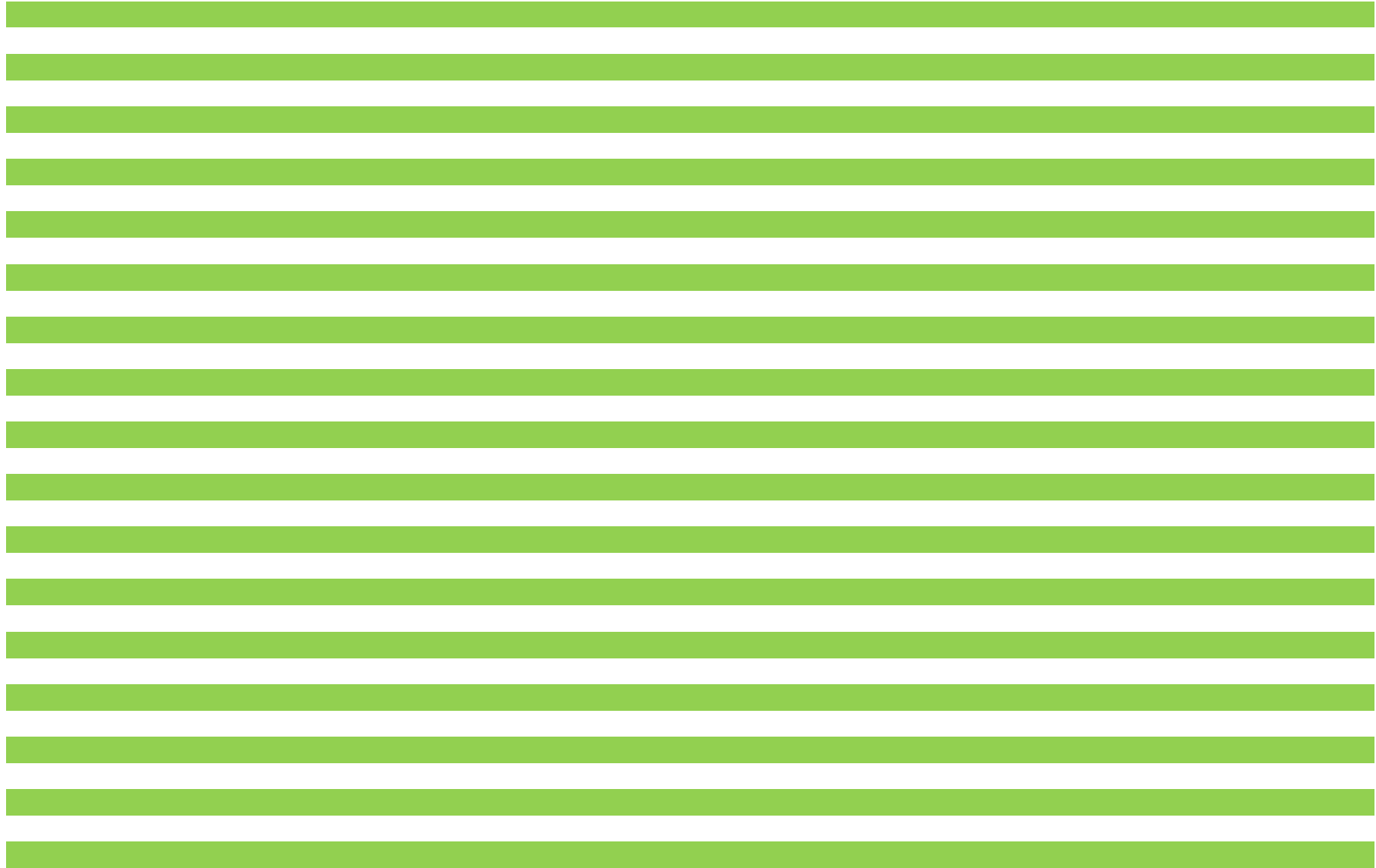
TAB]

TAB]

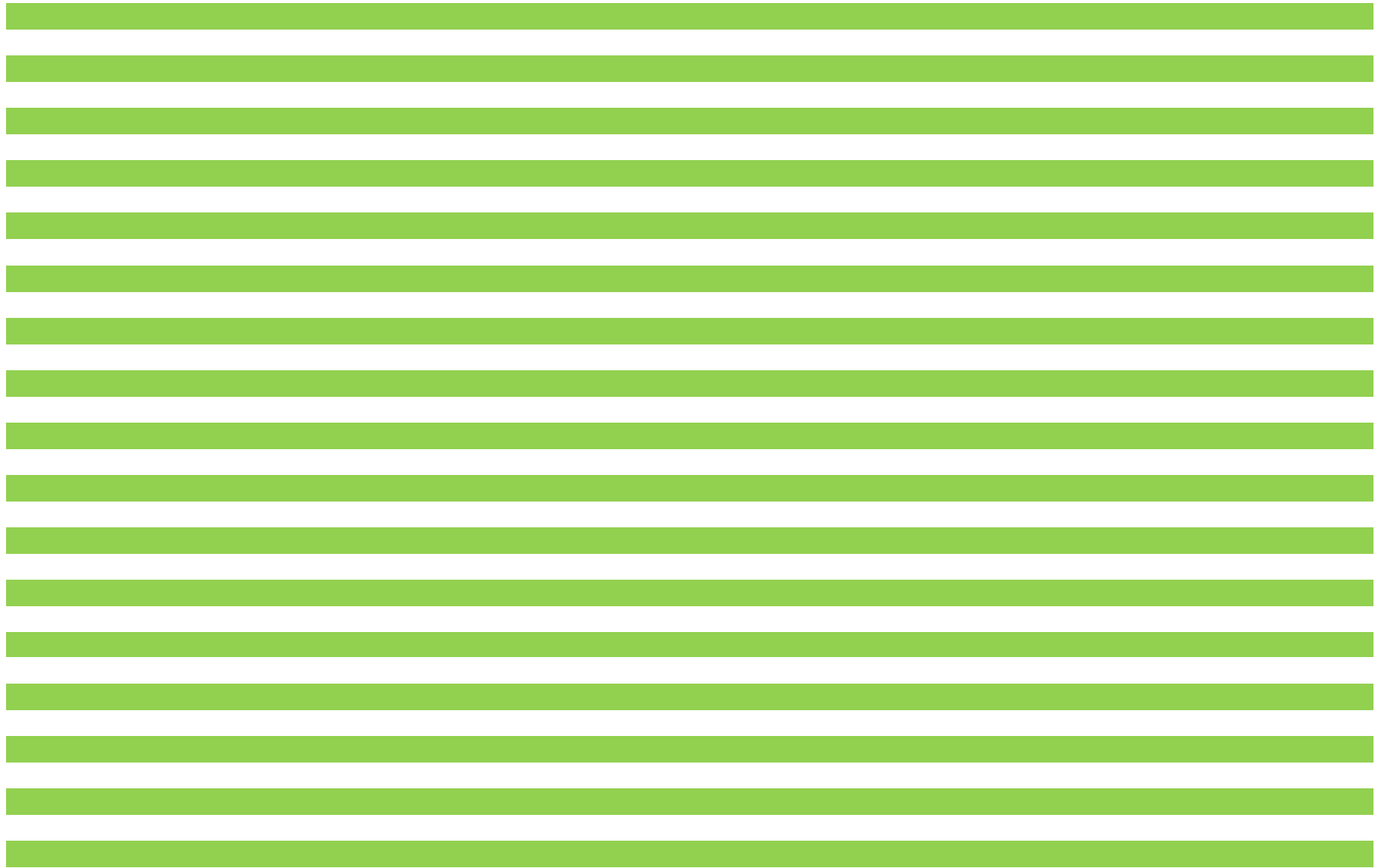
TAB]

TAB]

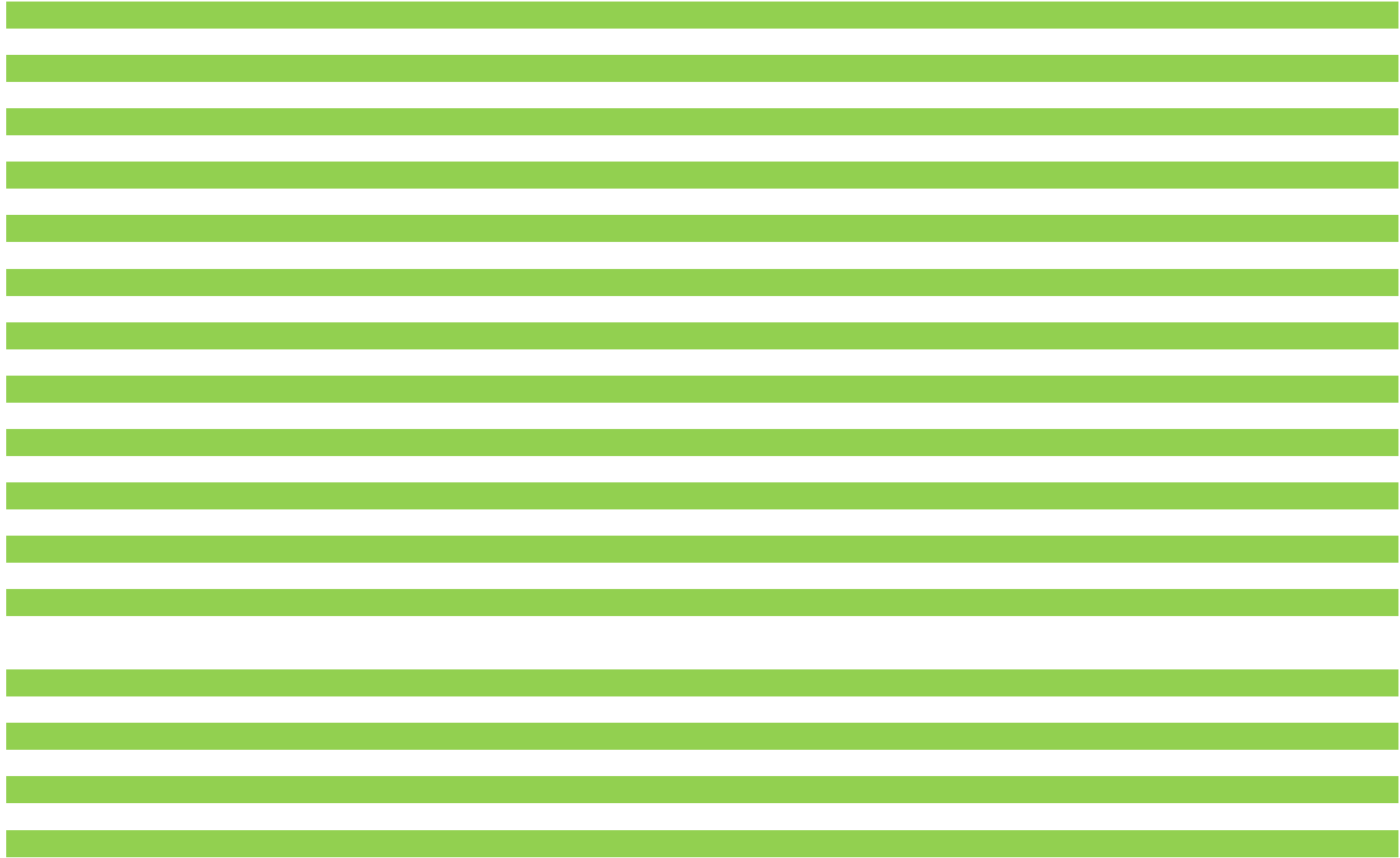
TAB]



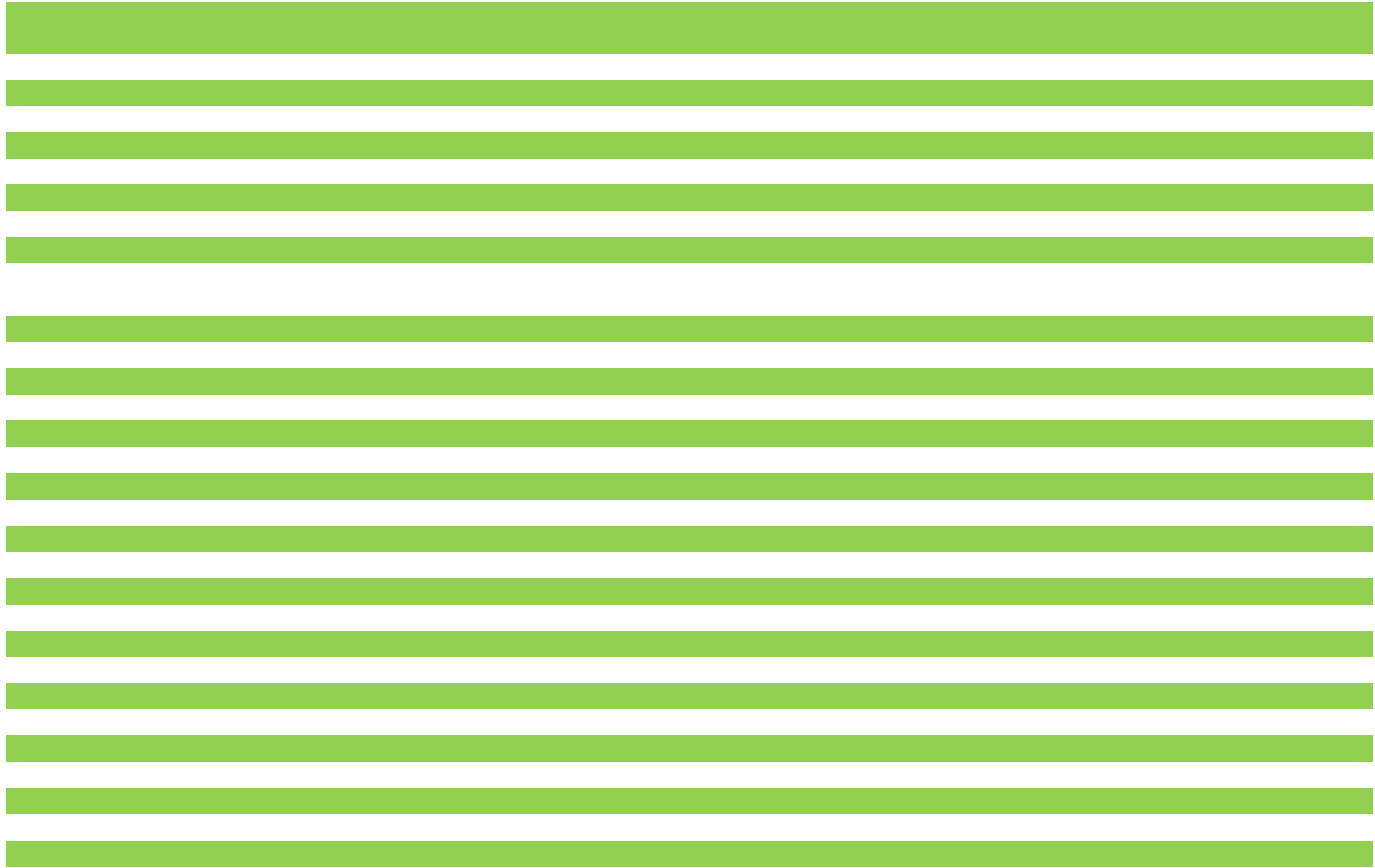
TAB]



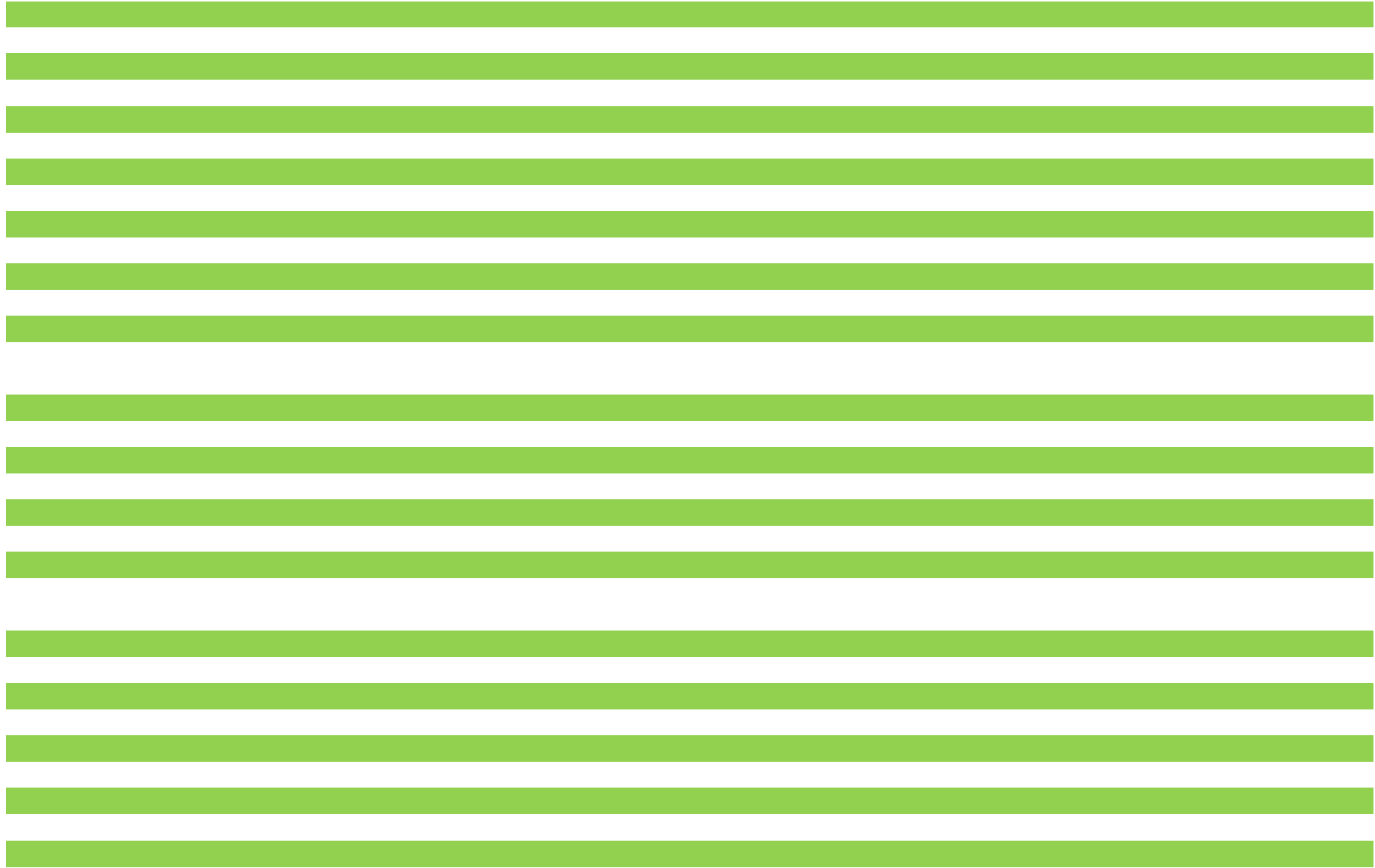
TAB]



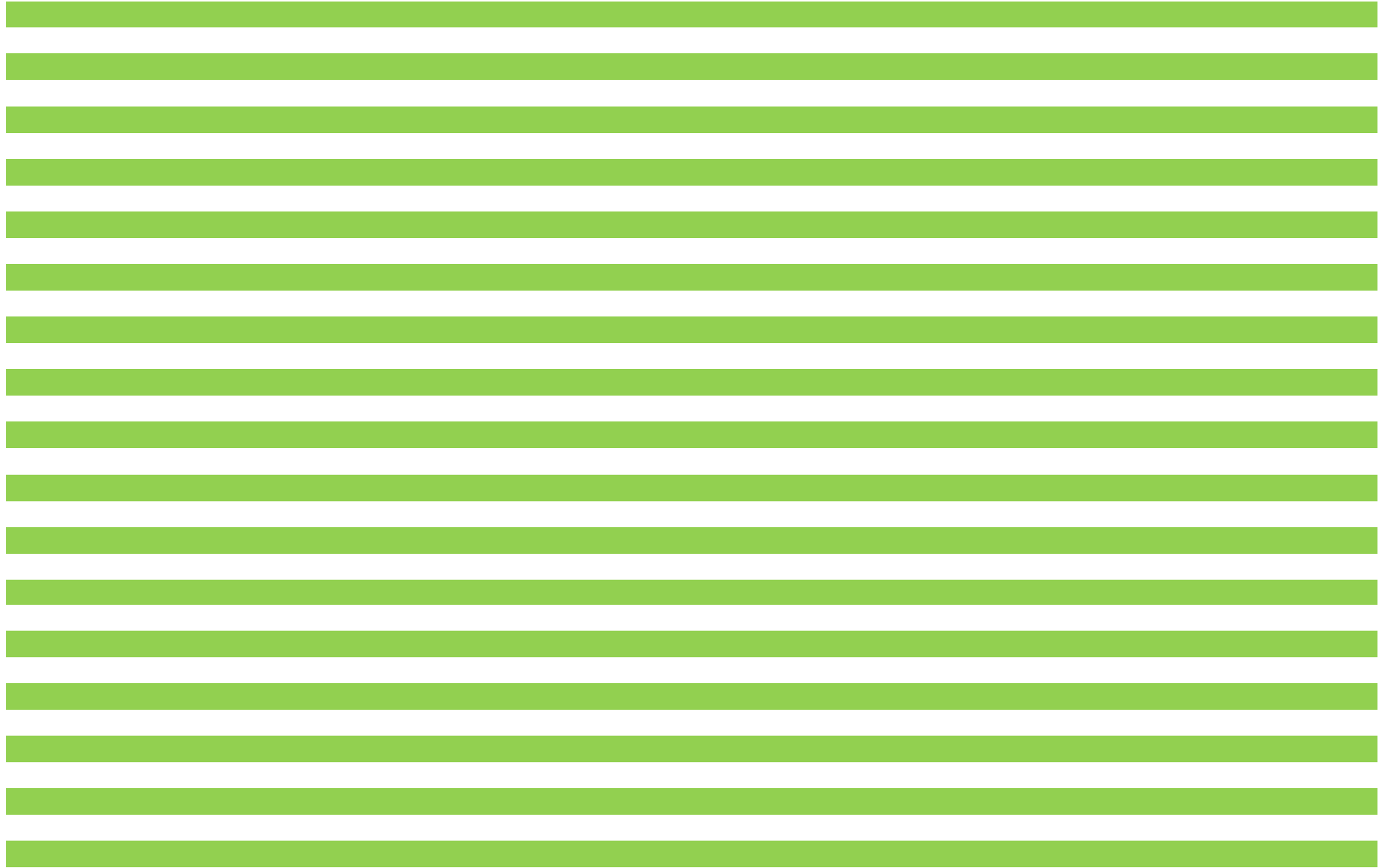
TAB]



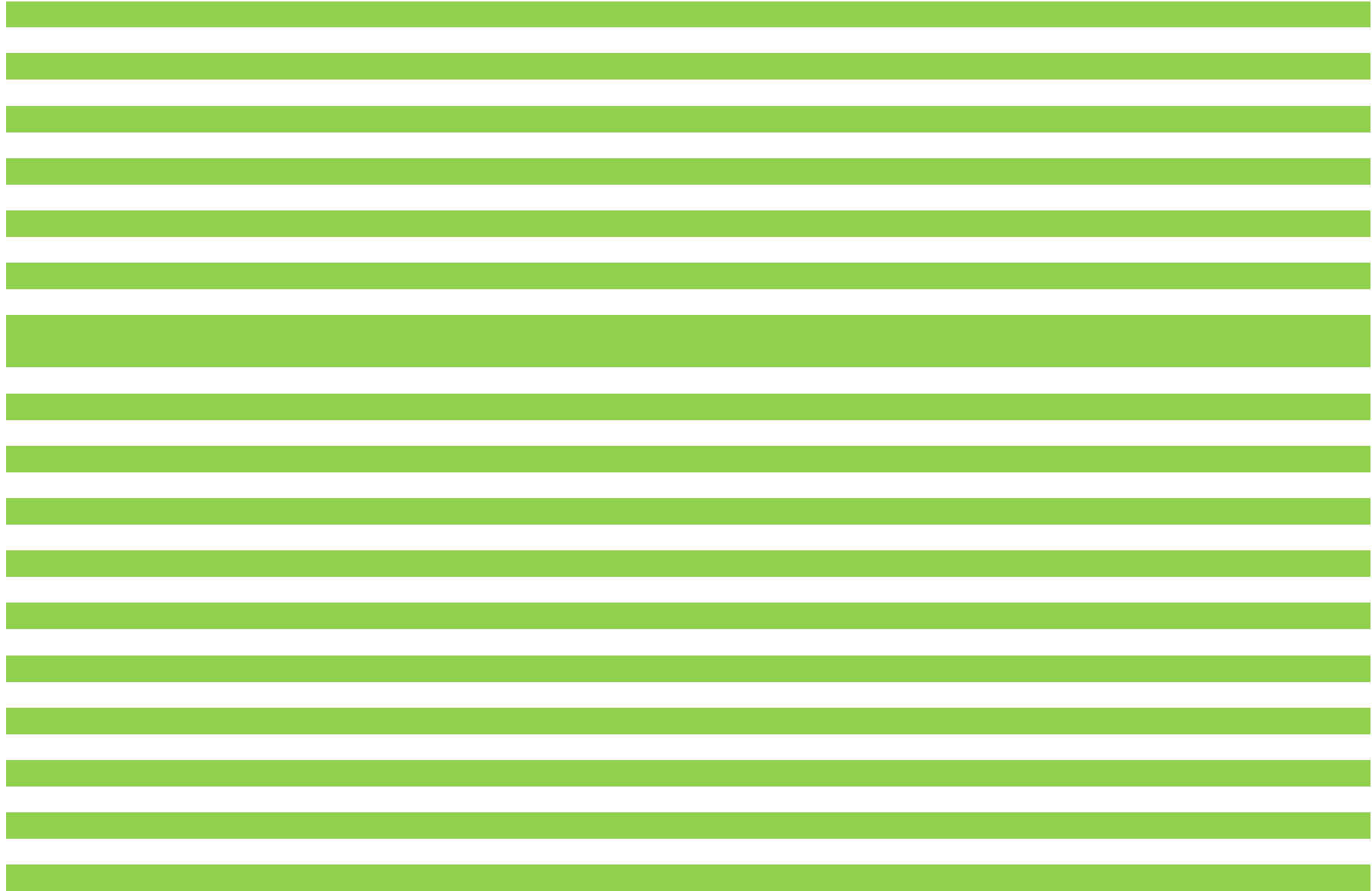
TAB]



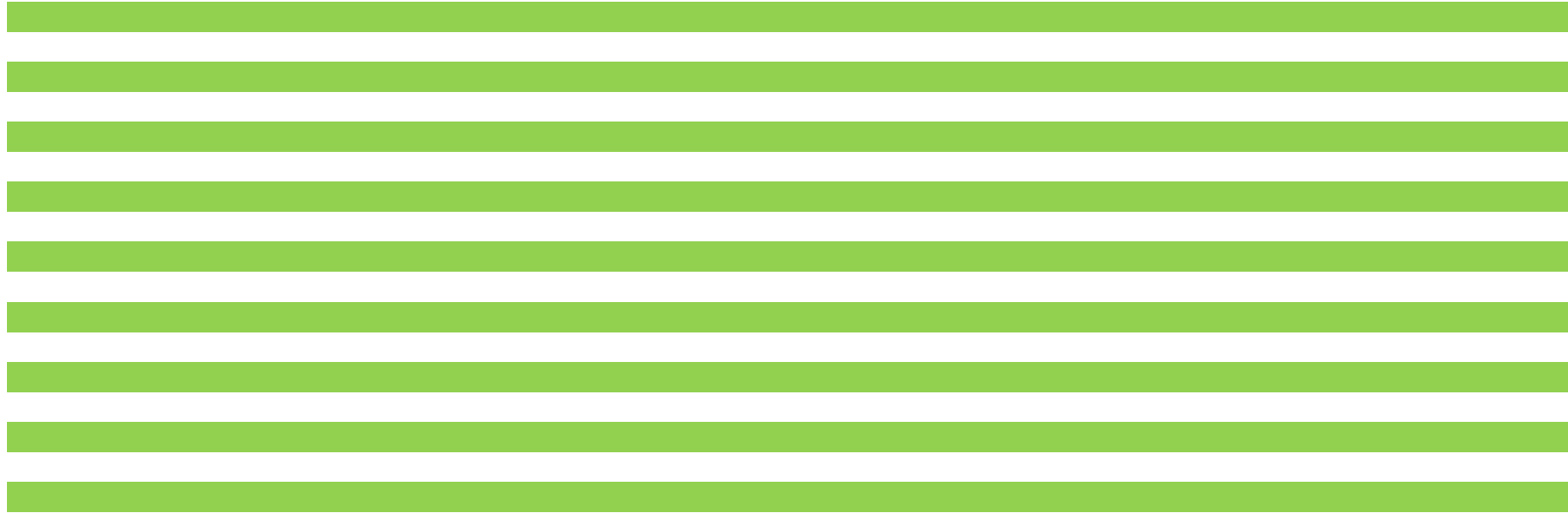
TAB]



TAB]



TAB]



TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

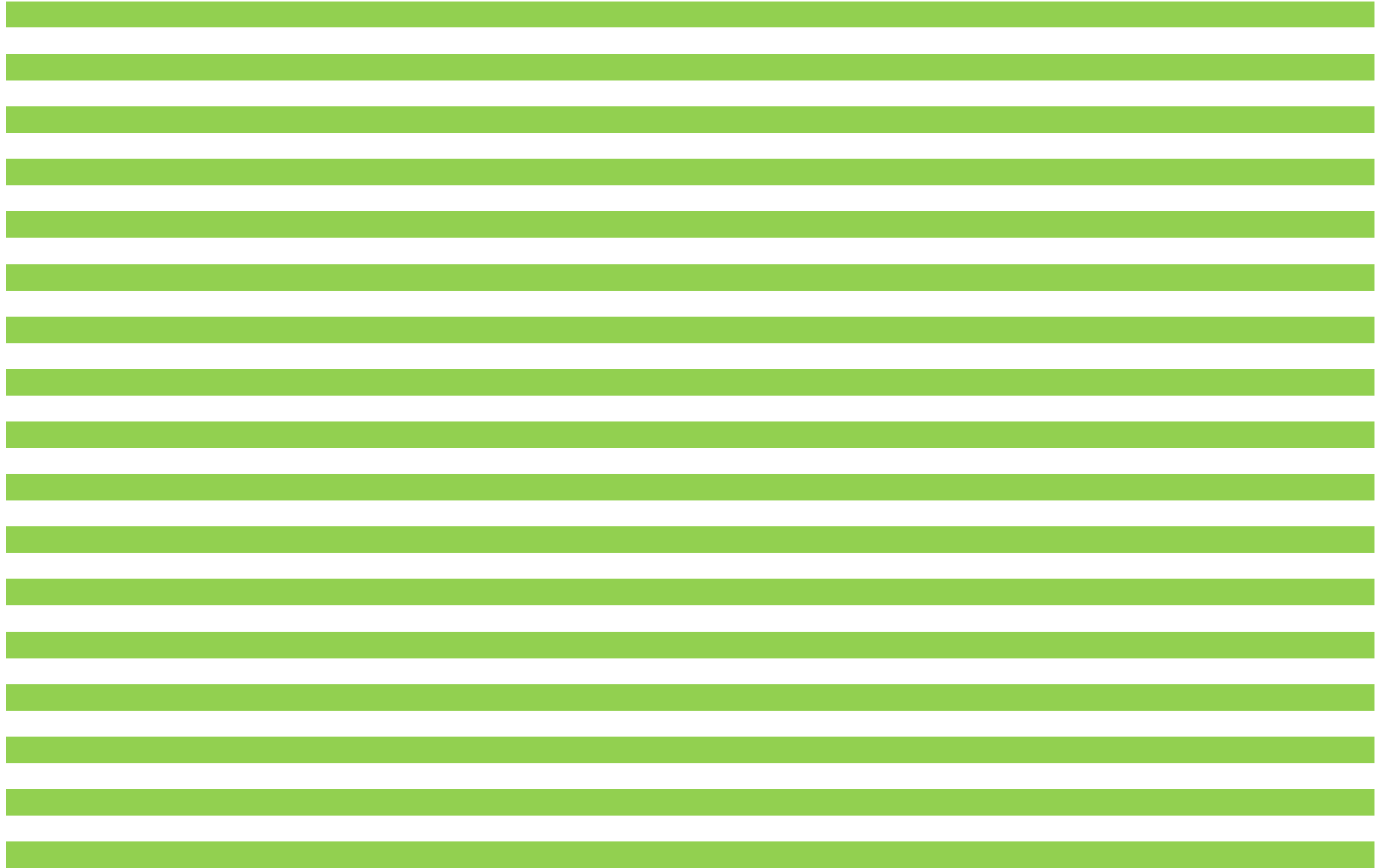
TAB]

TAB]

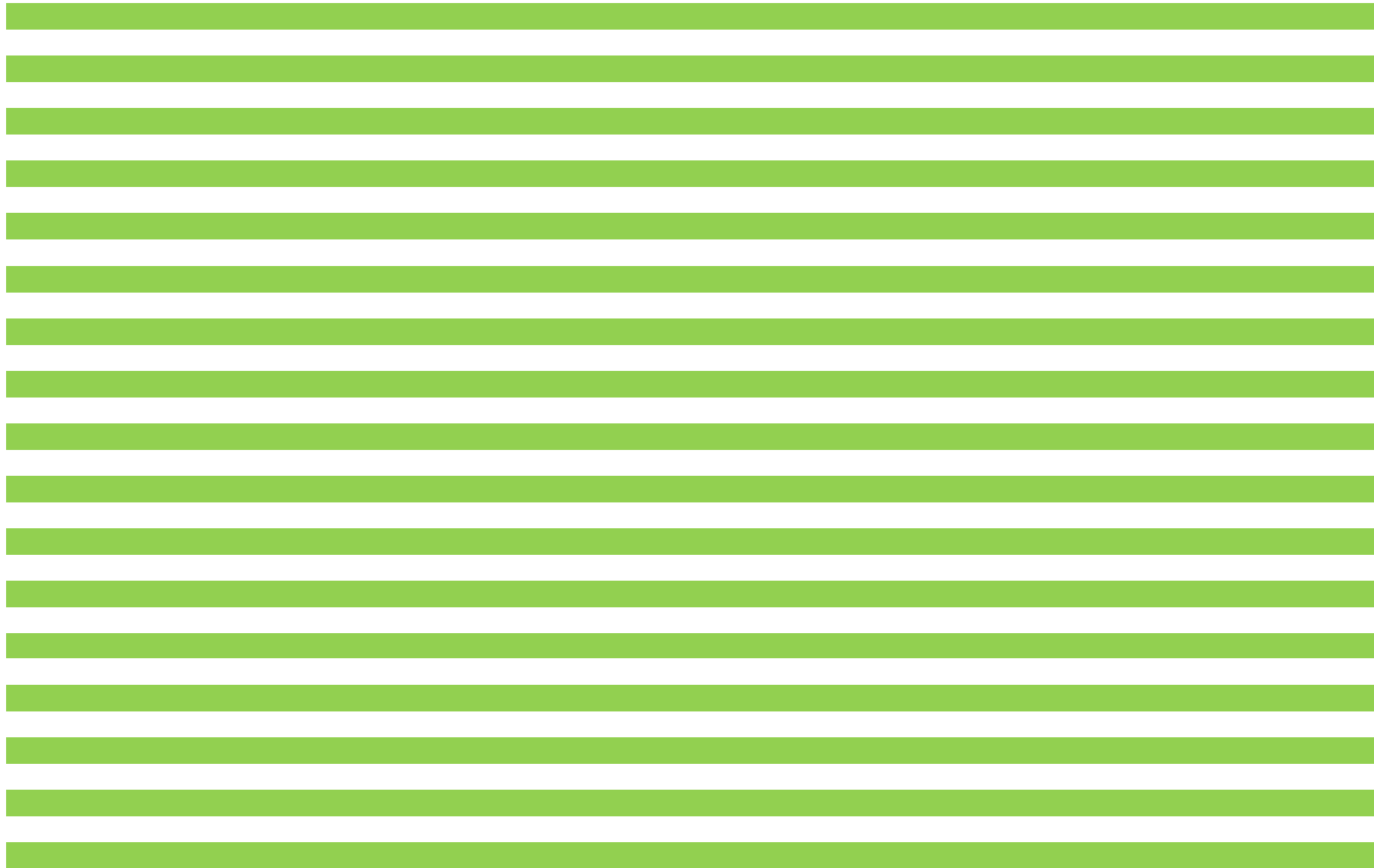
TAB]

TAB]

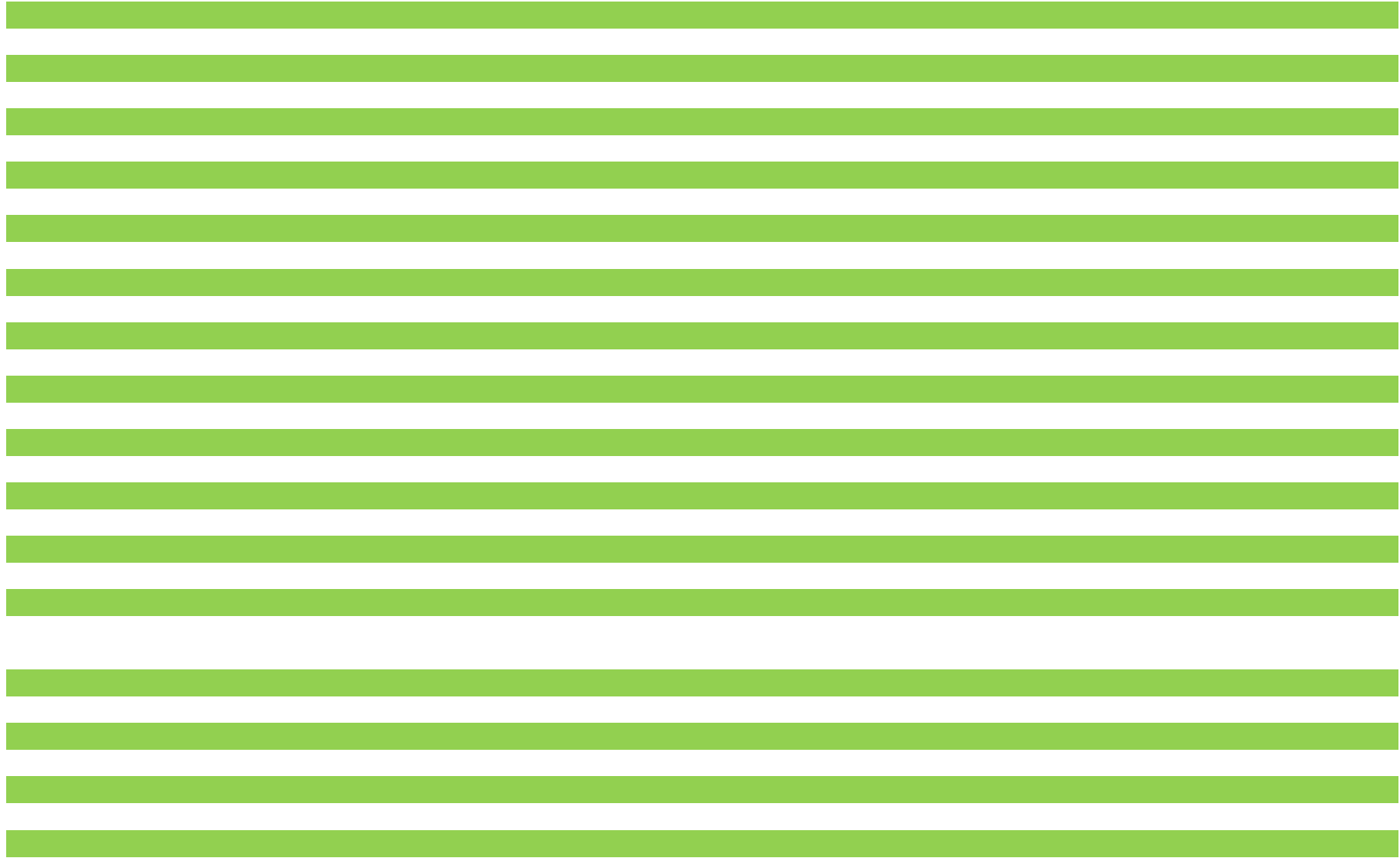
TAB]



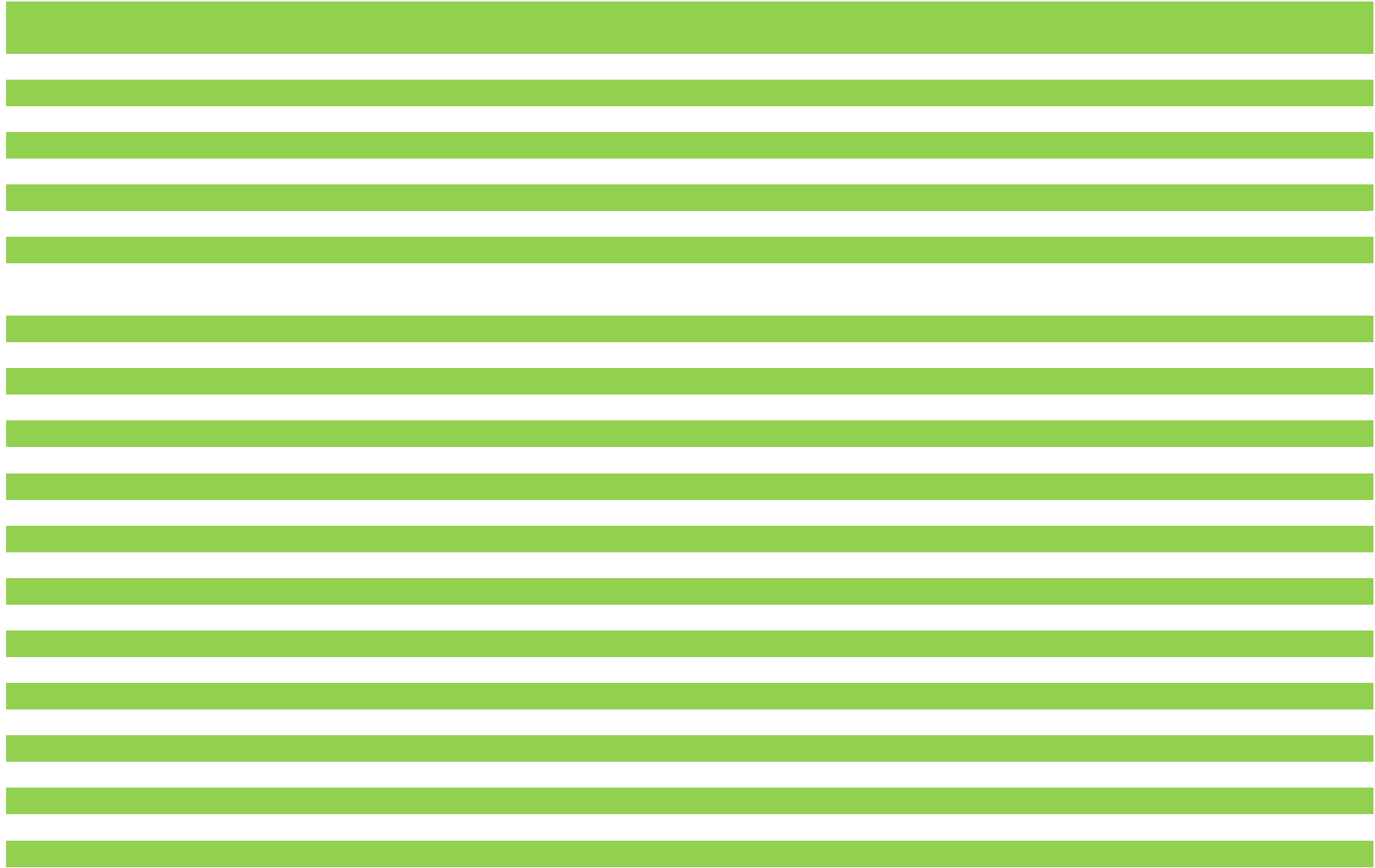
TAB]



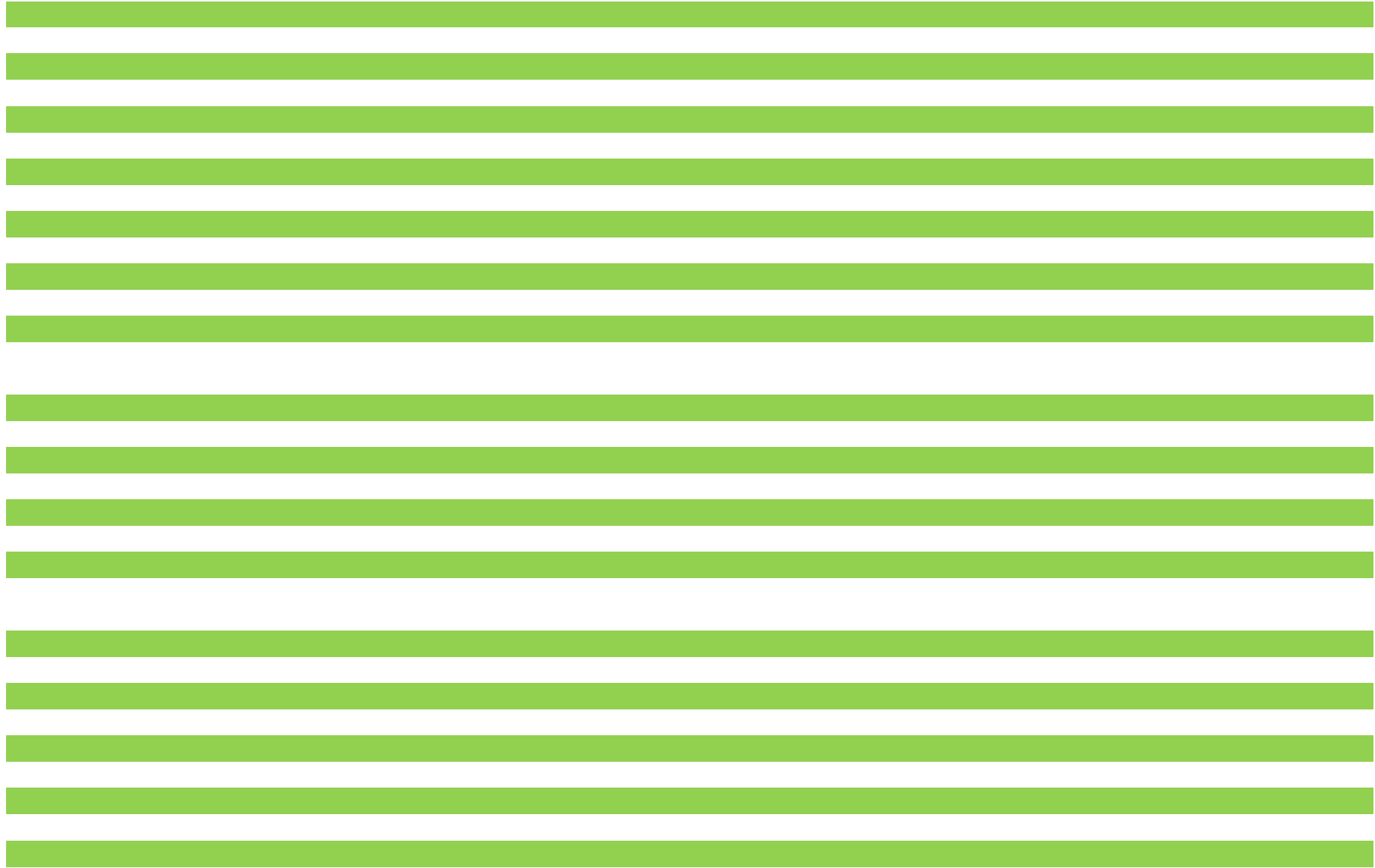
TAB]



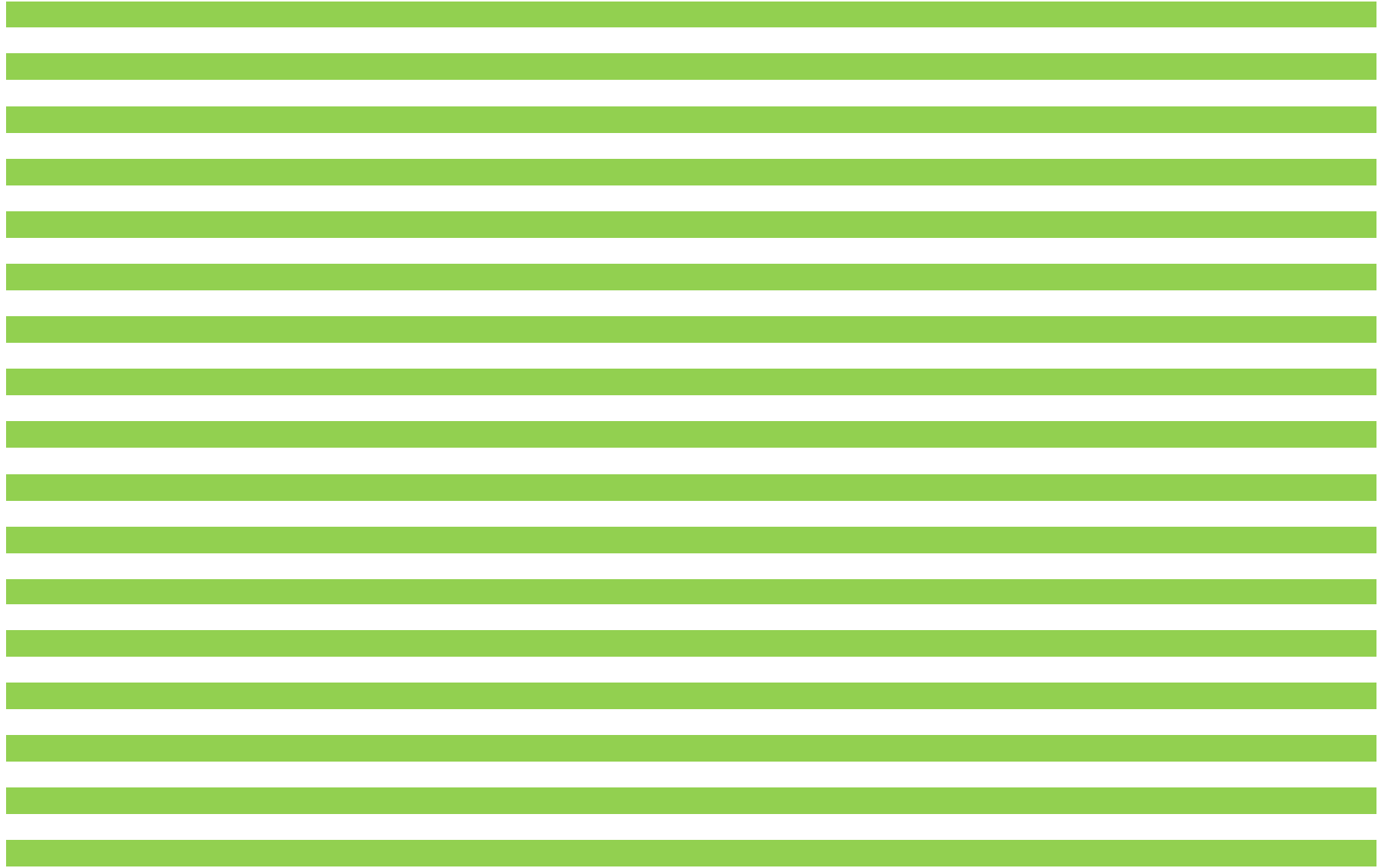
TAB]



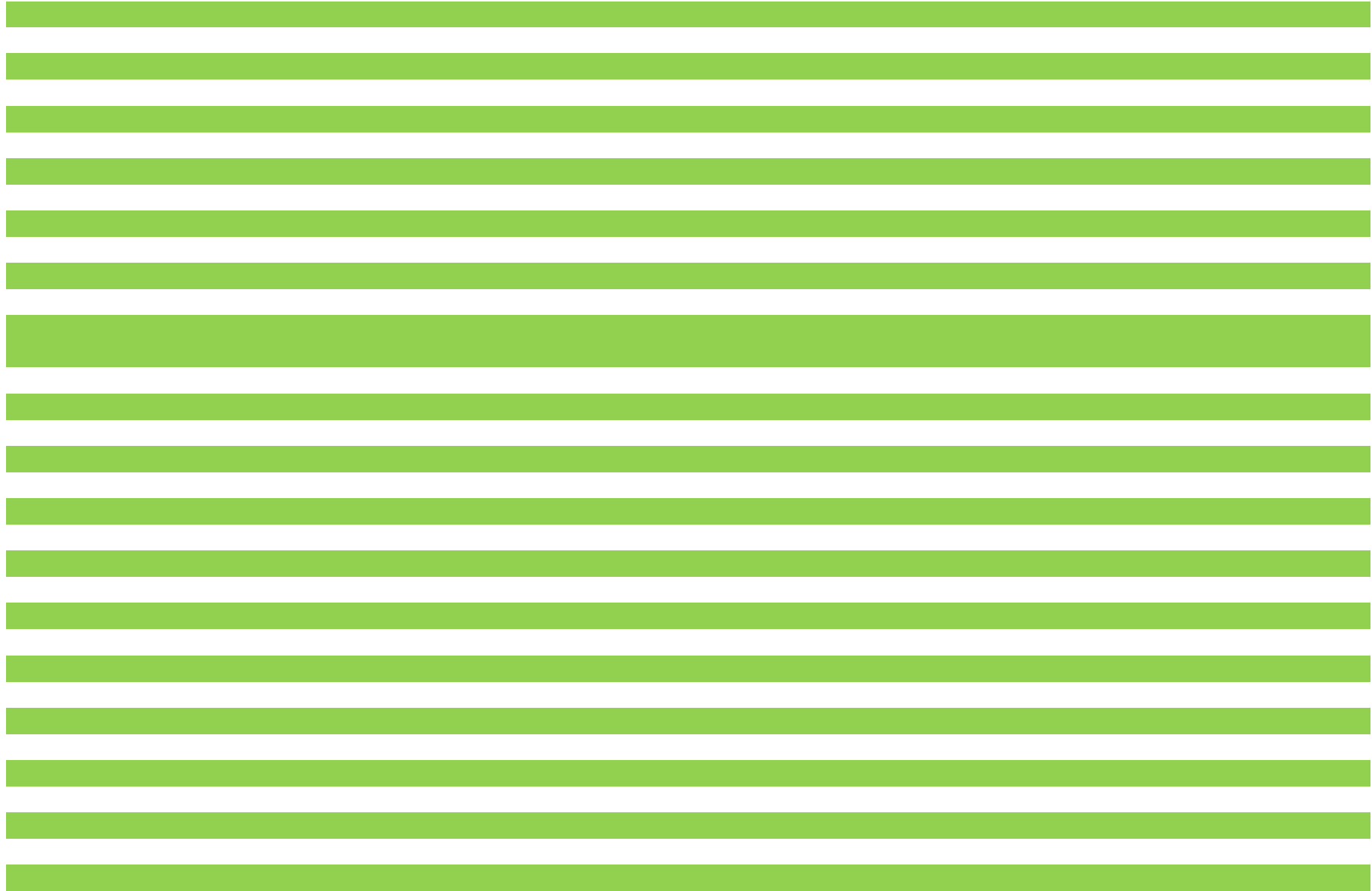
TAB]



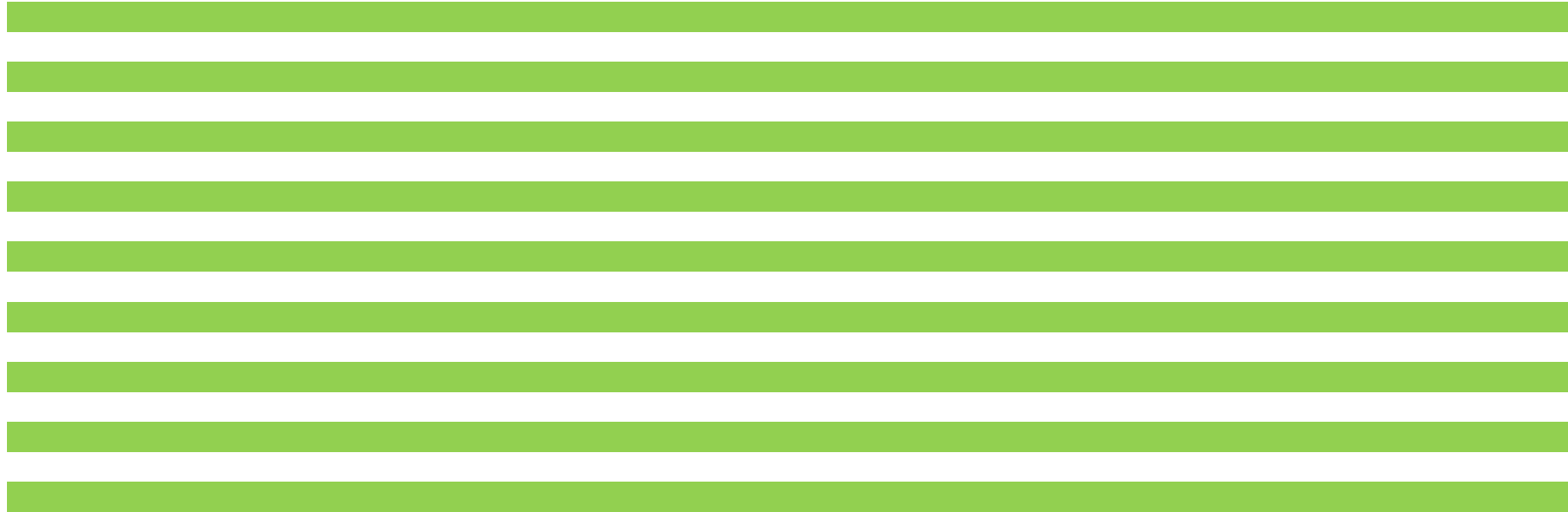
TAB]



TAB]



TAB]



TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

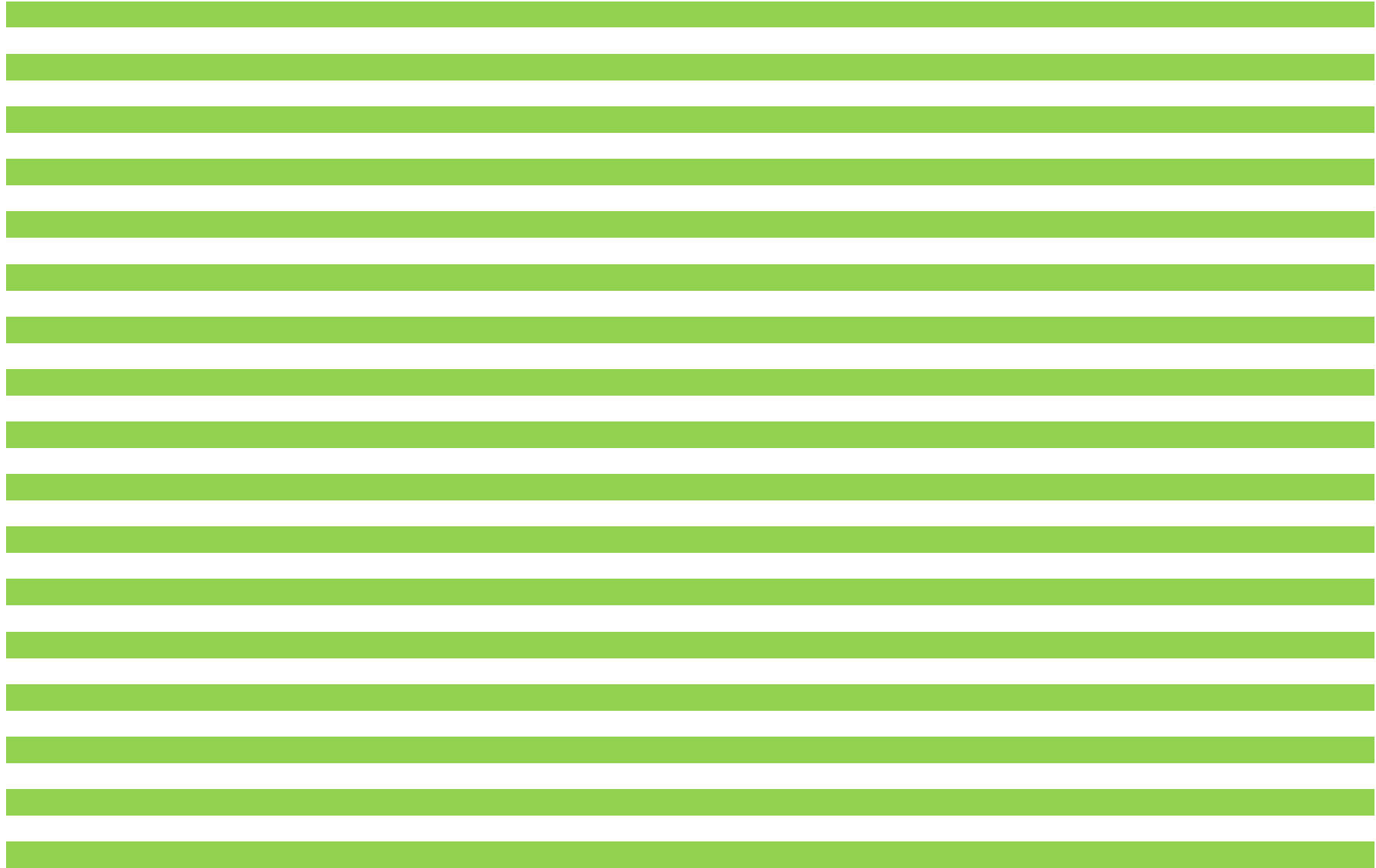
TAB]

TAB]

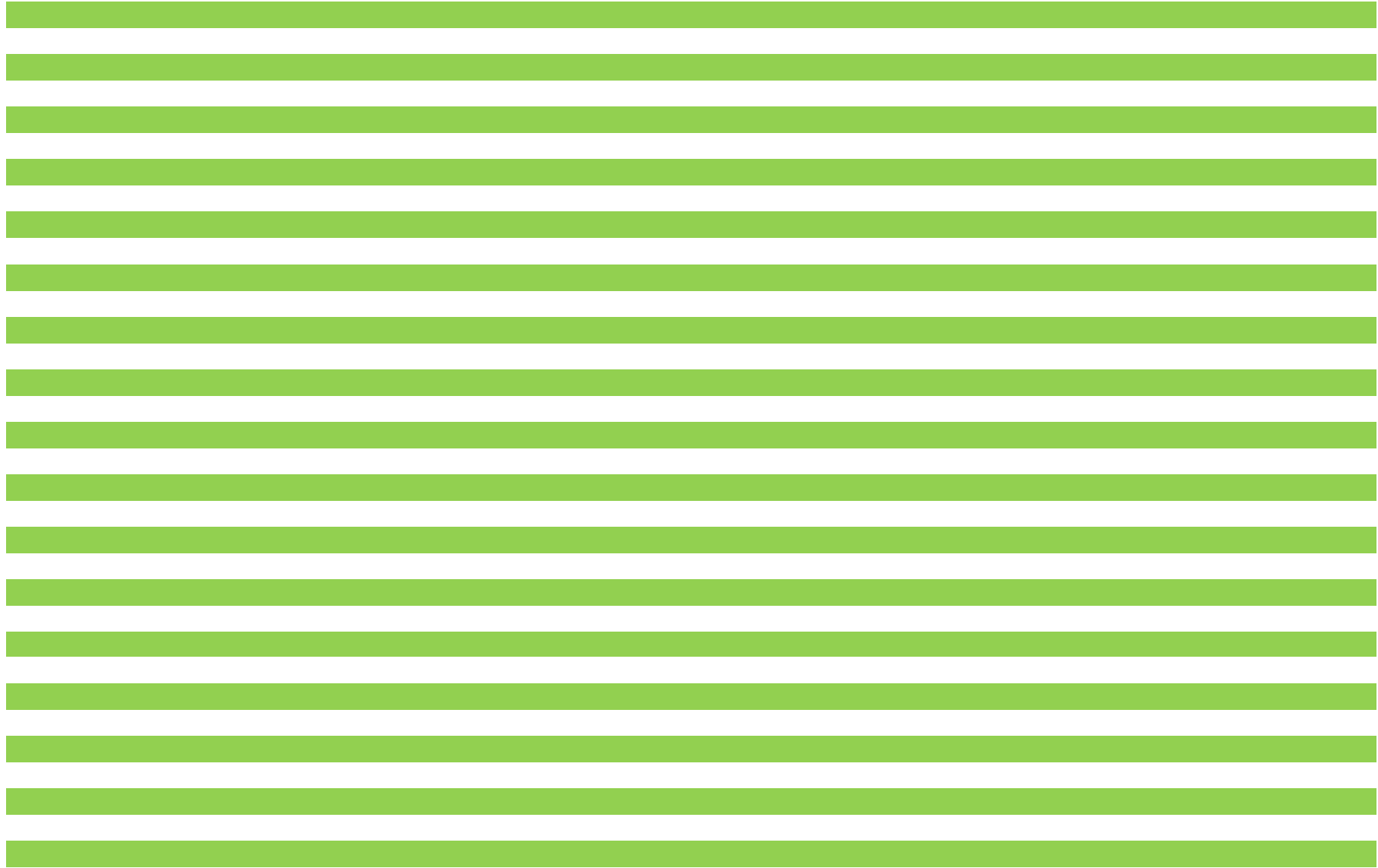
TAB]

TAB]

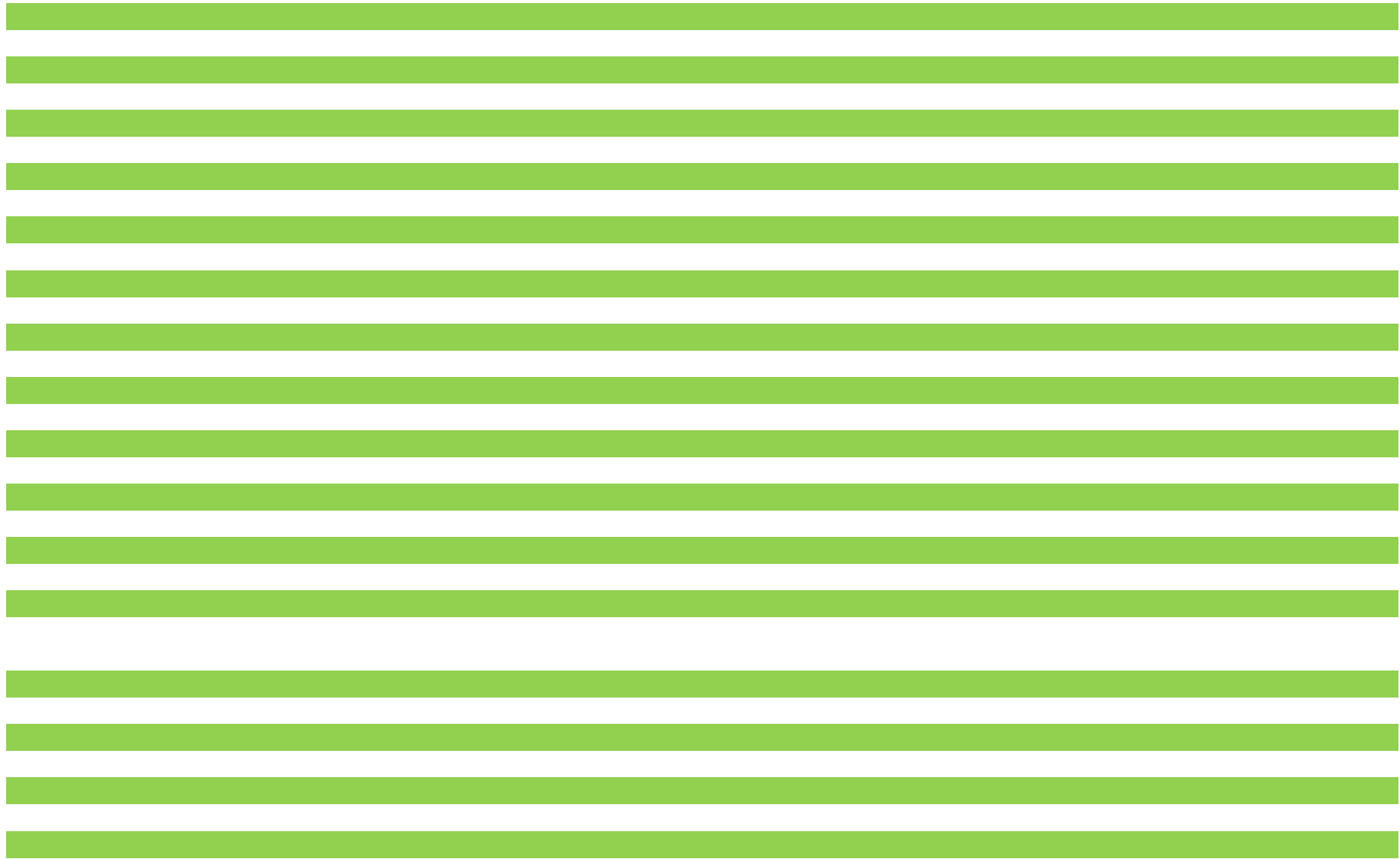
TAB]



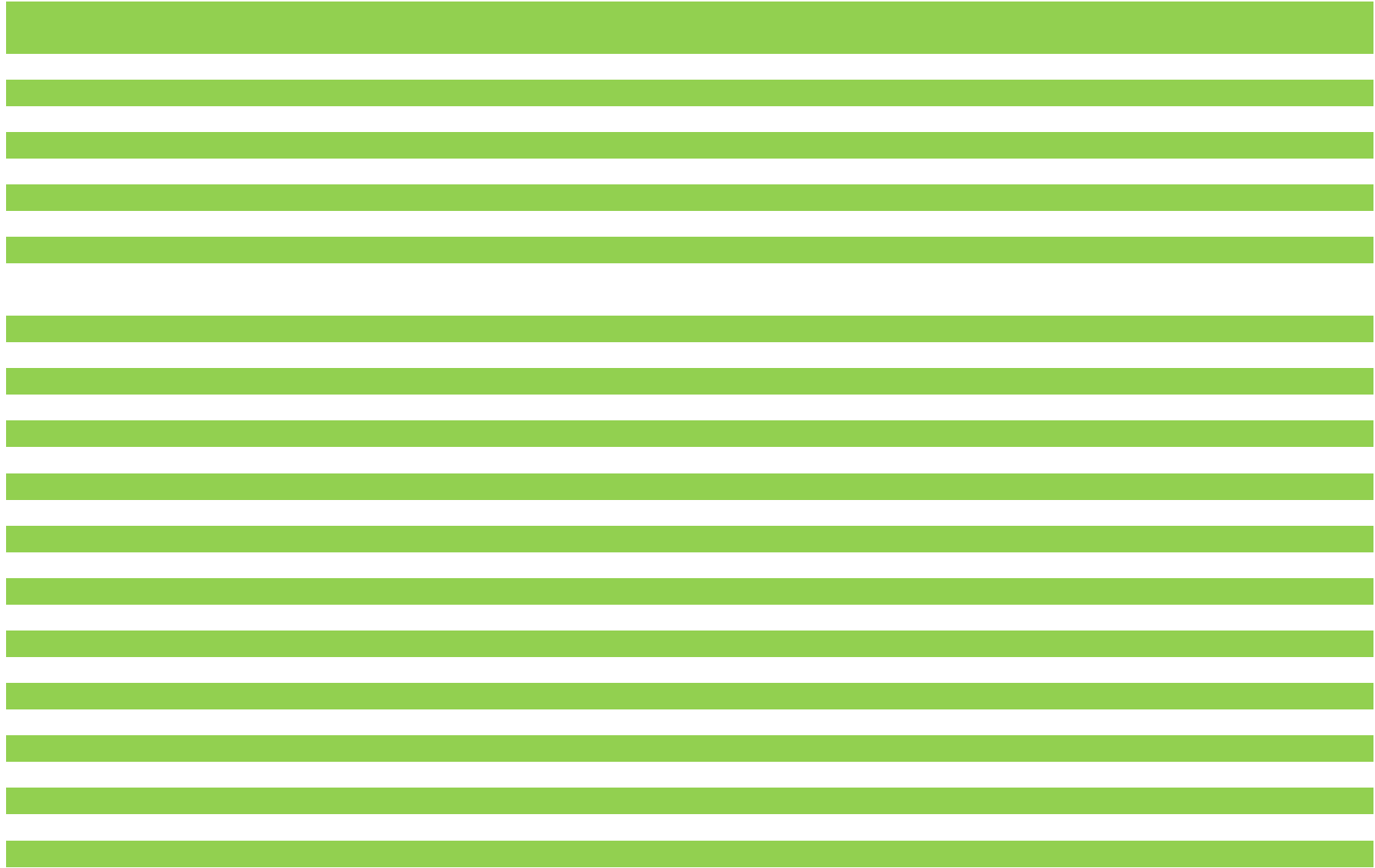
TAB]



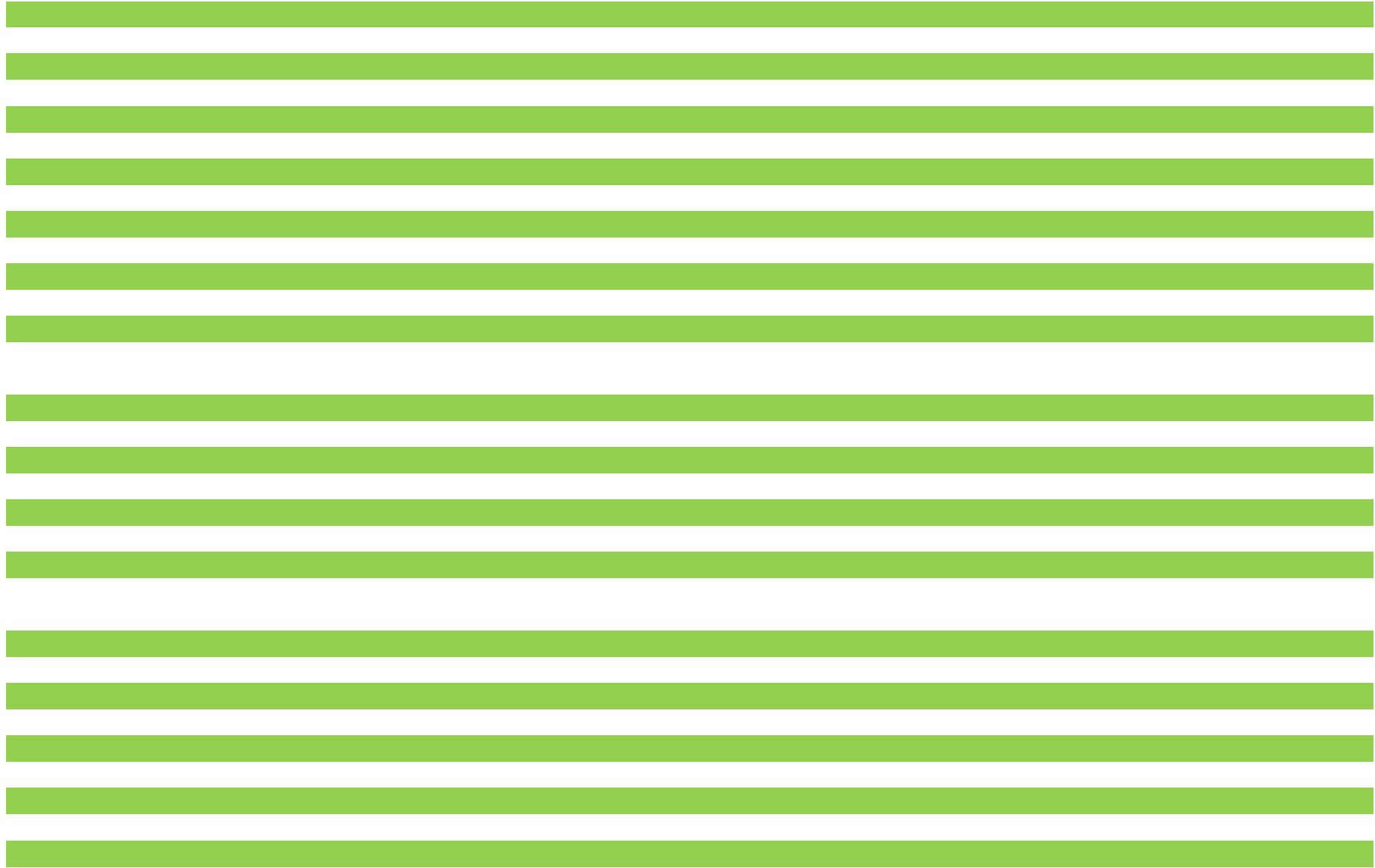
TAB]



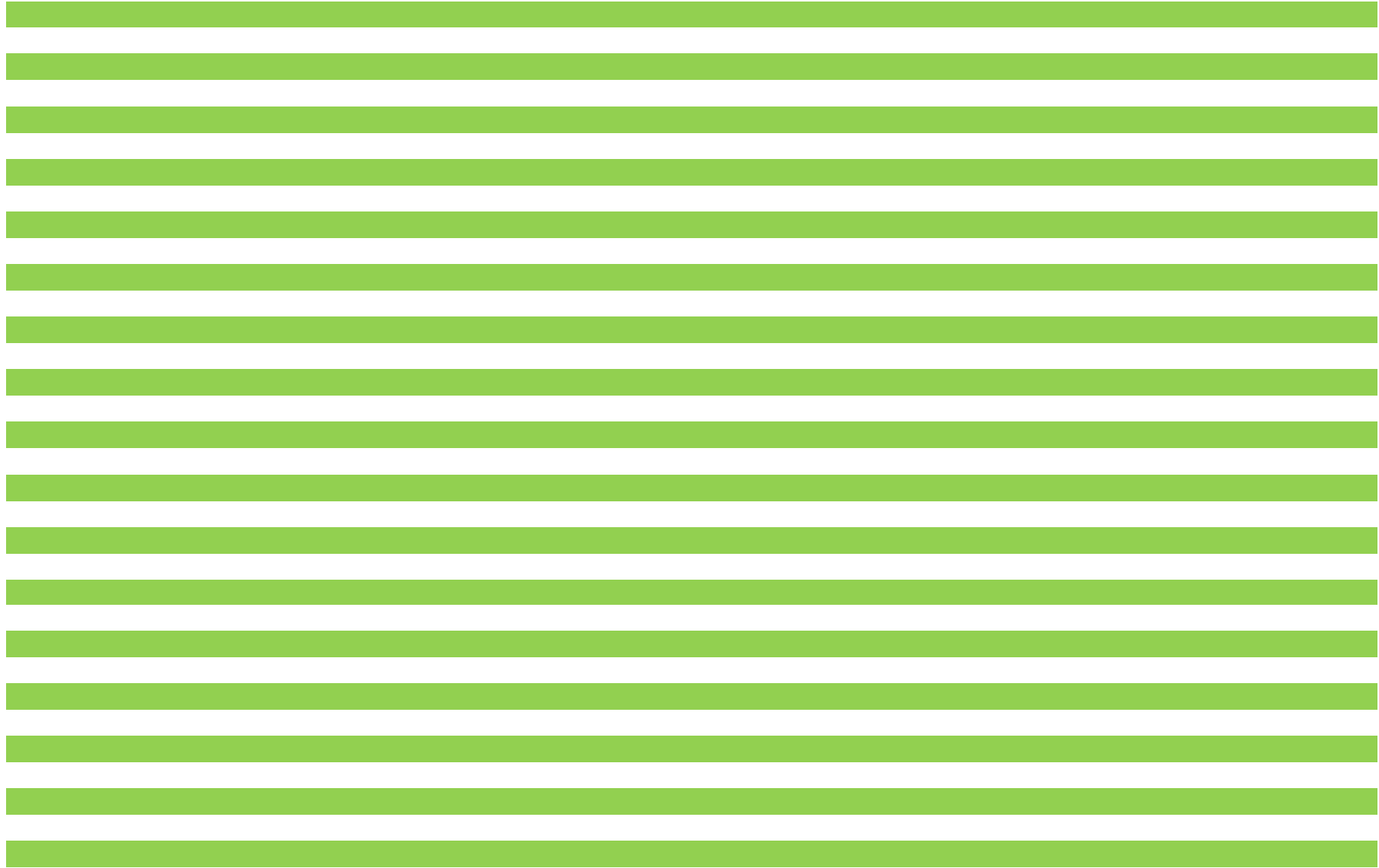
TAB]



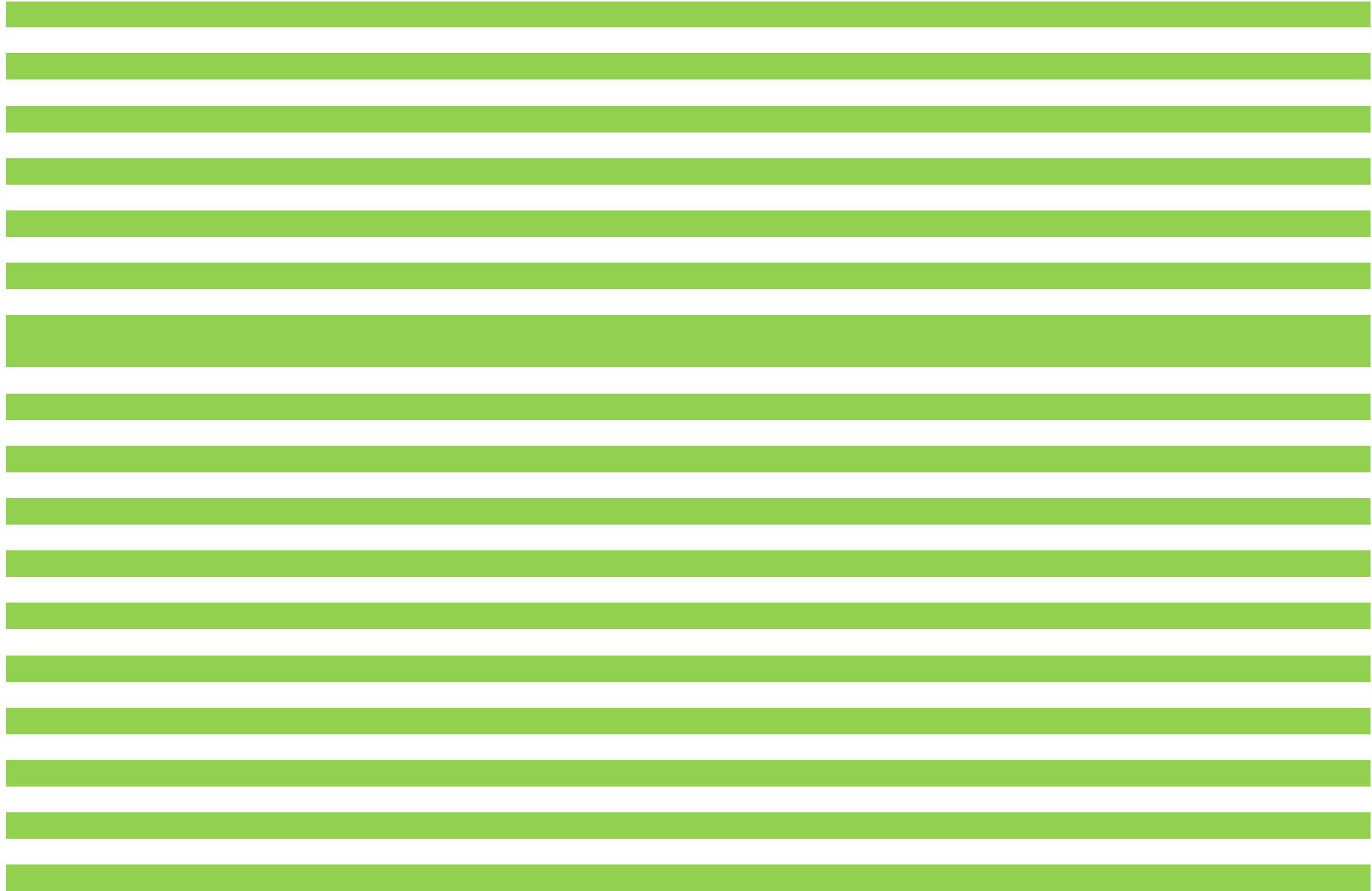
TAB]



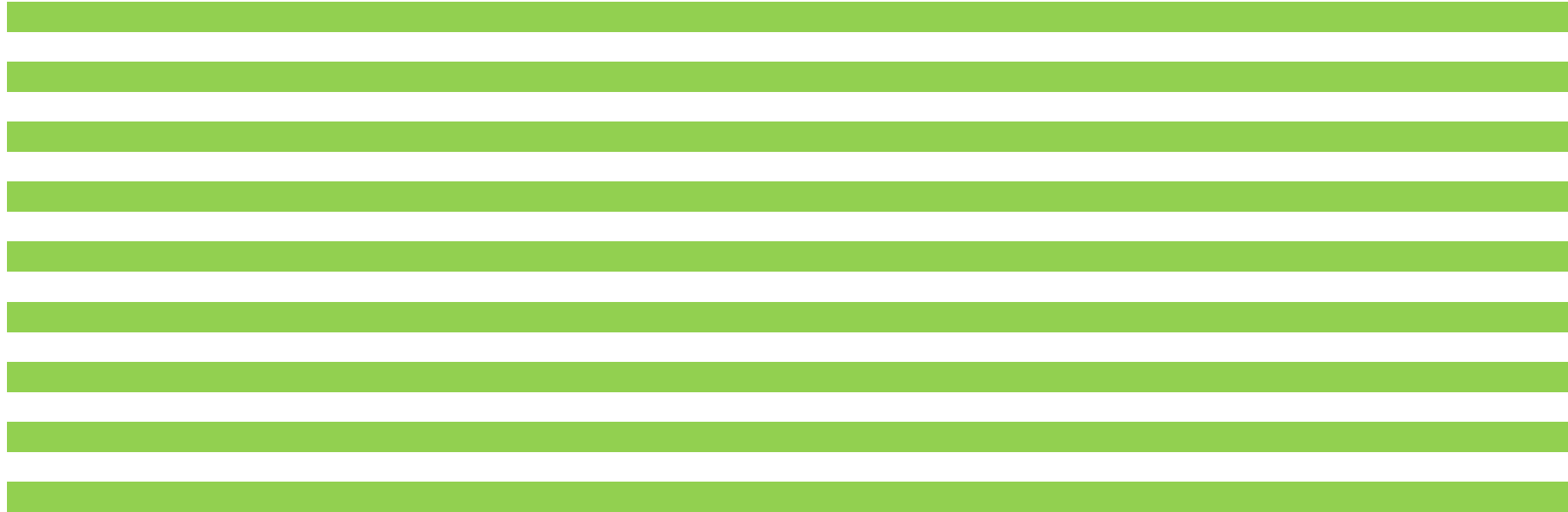
TAB]



TAB]



TAB]



TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

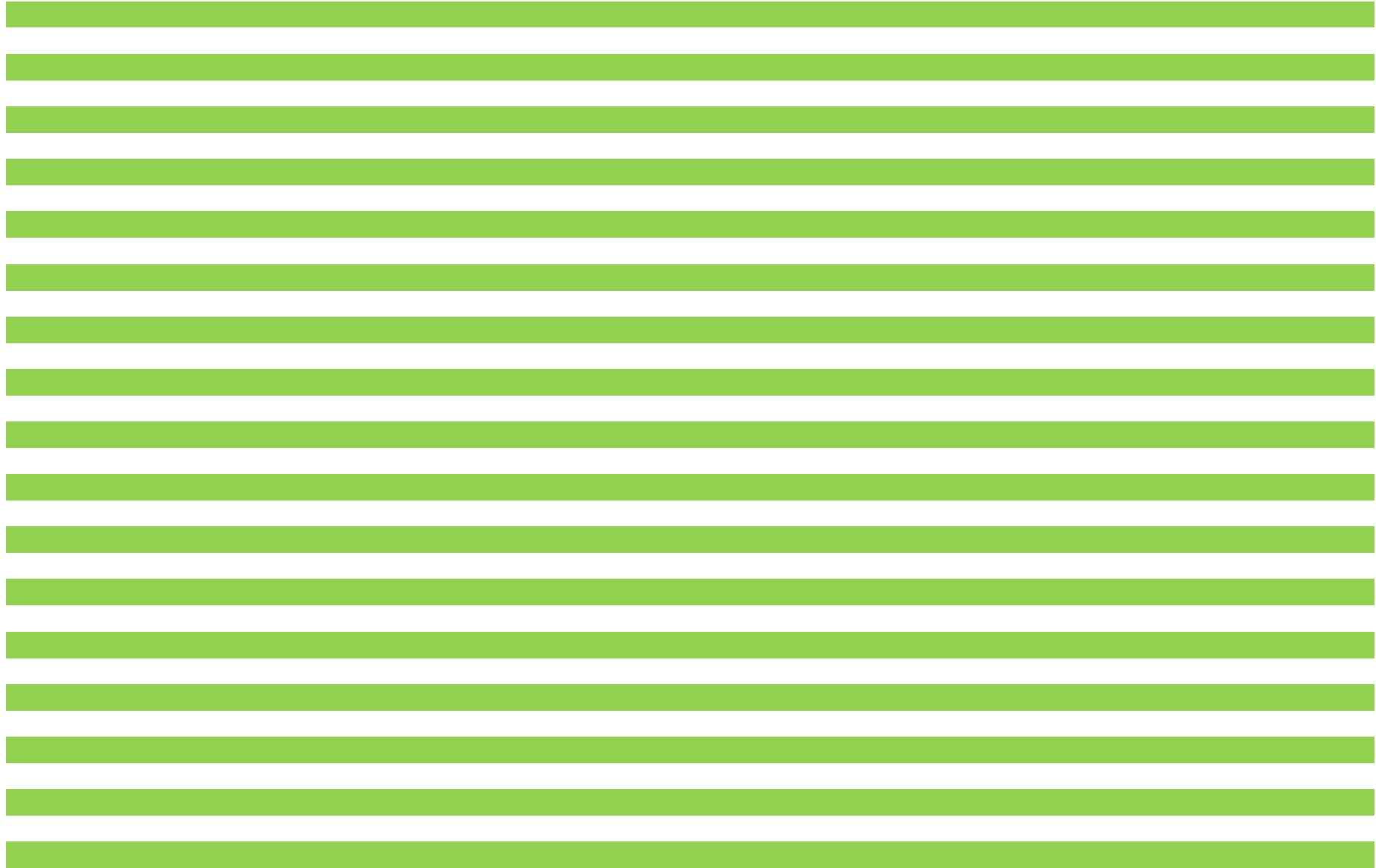
TAB]

TAB]

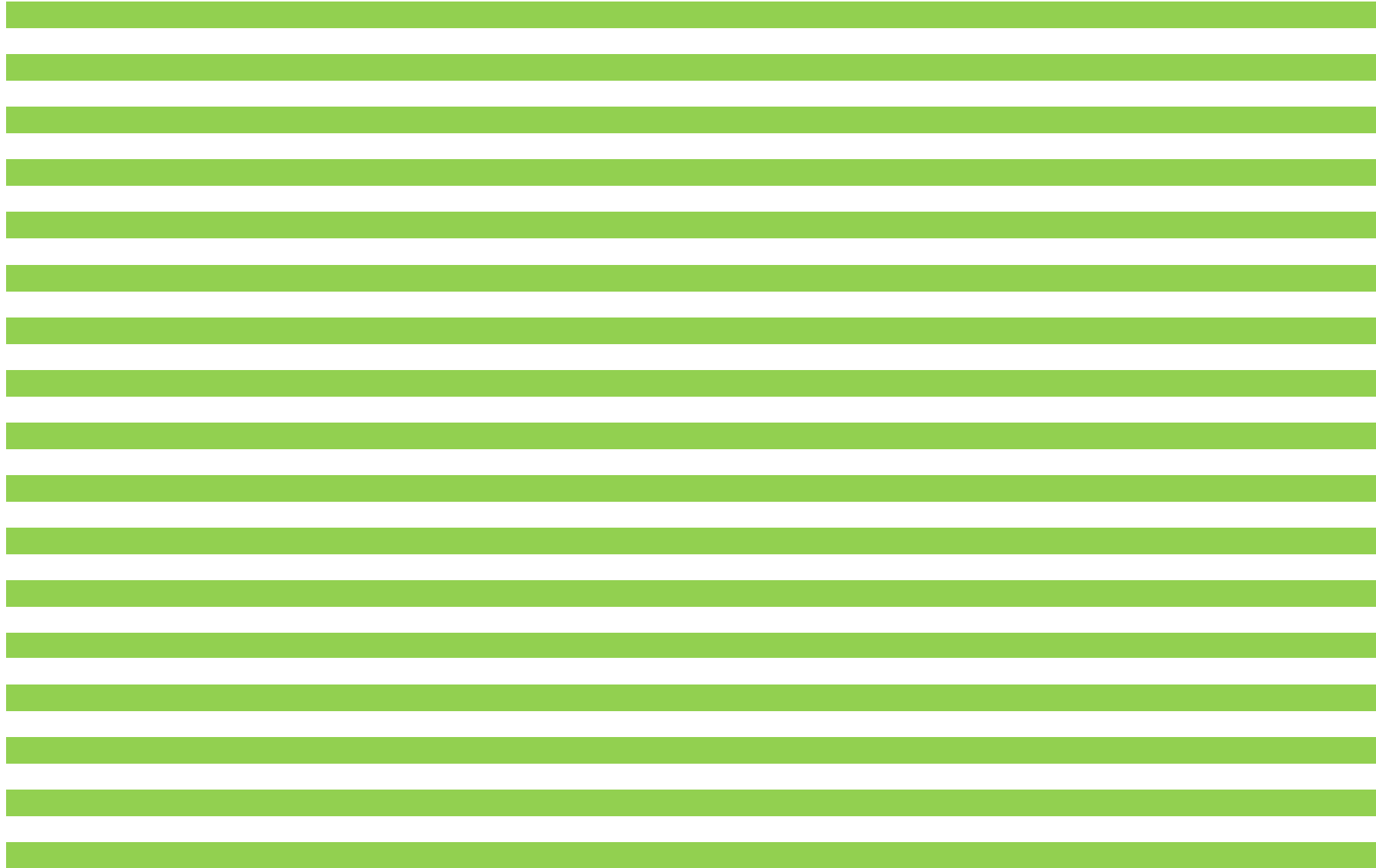
TAB]

TAB]

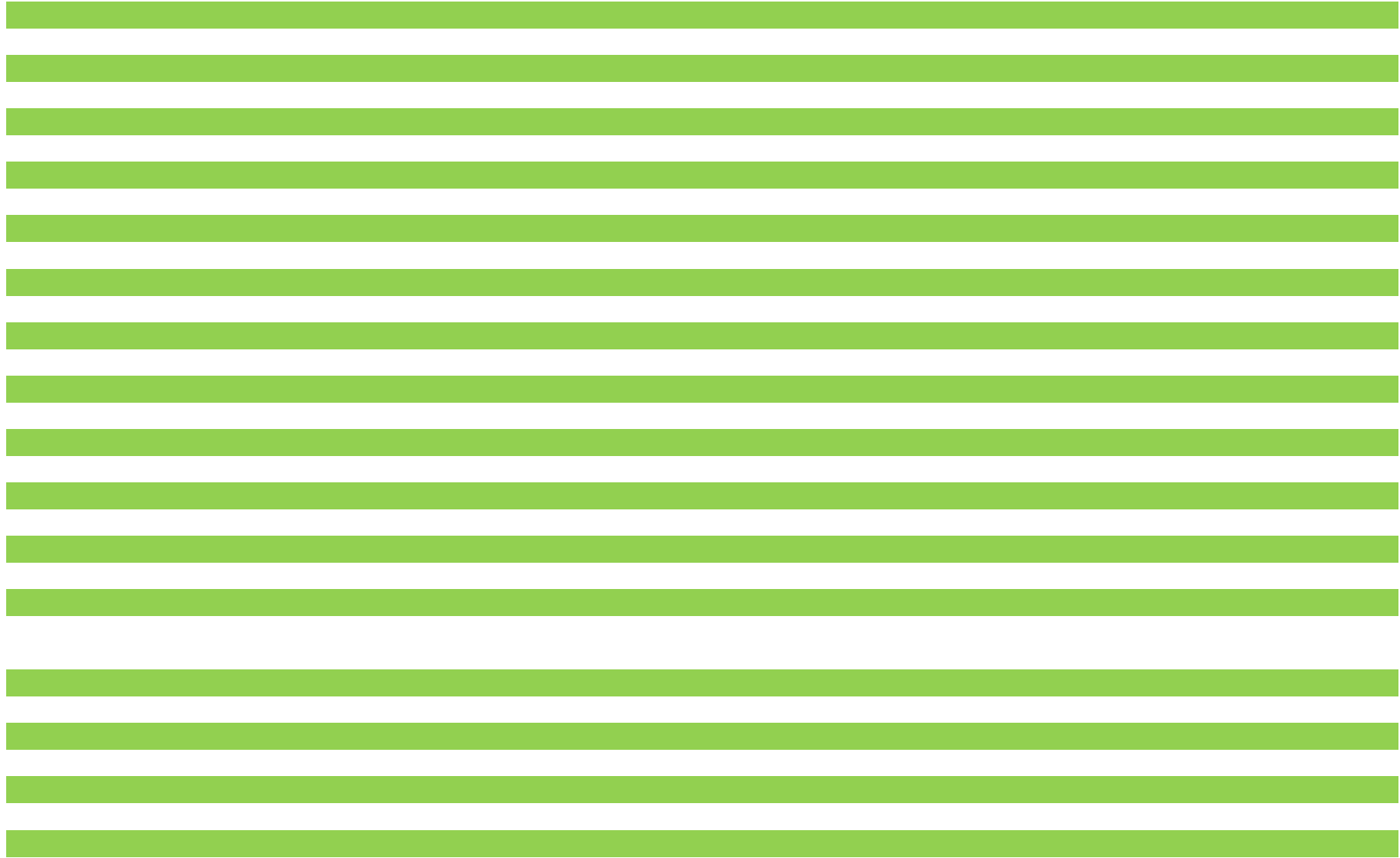
TAB]



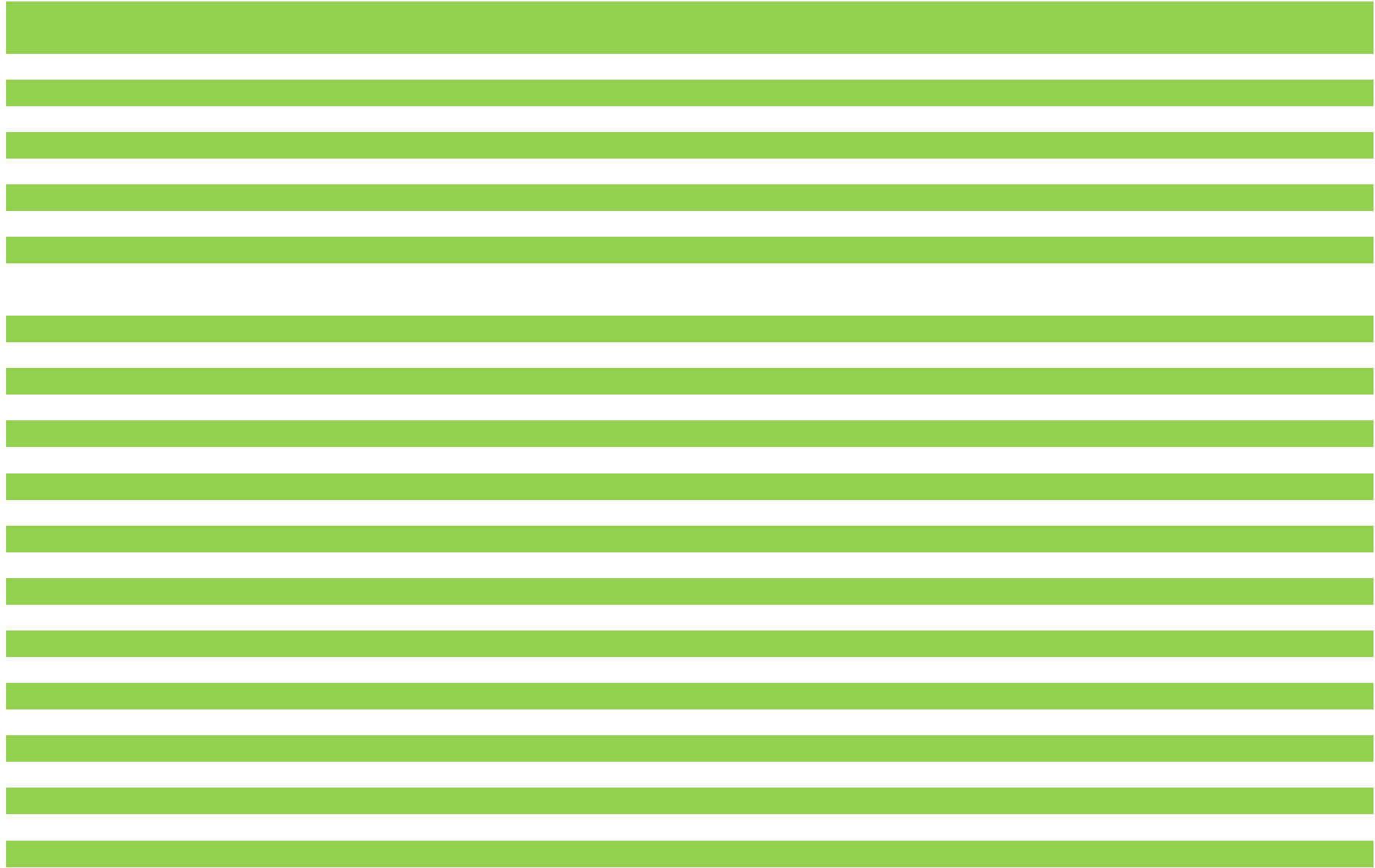
TAB]



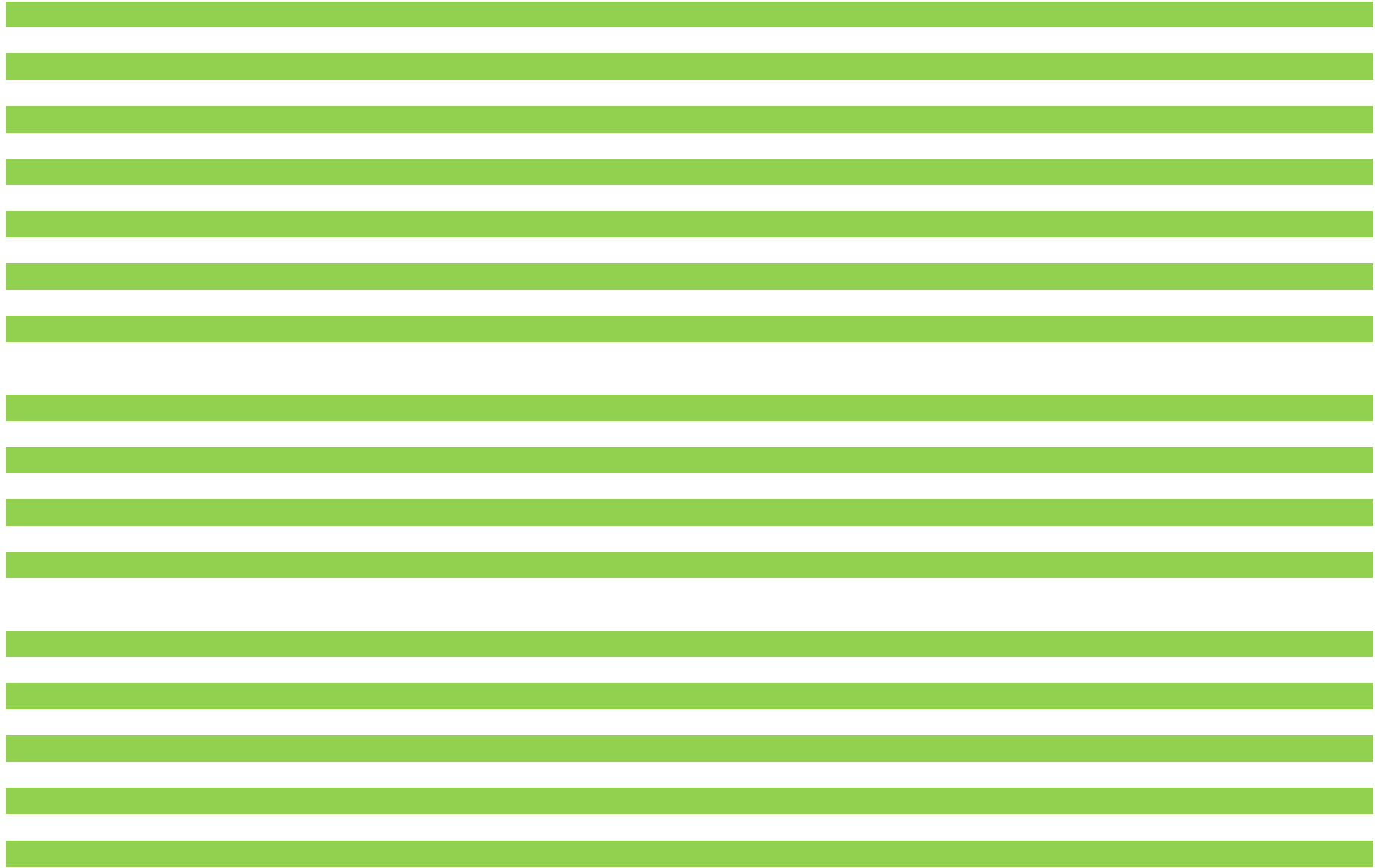
TAB]



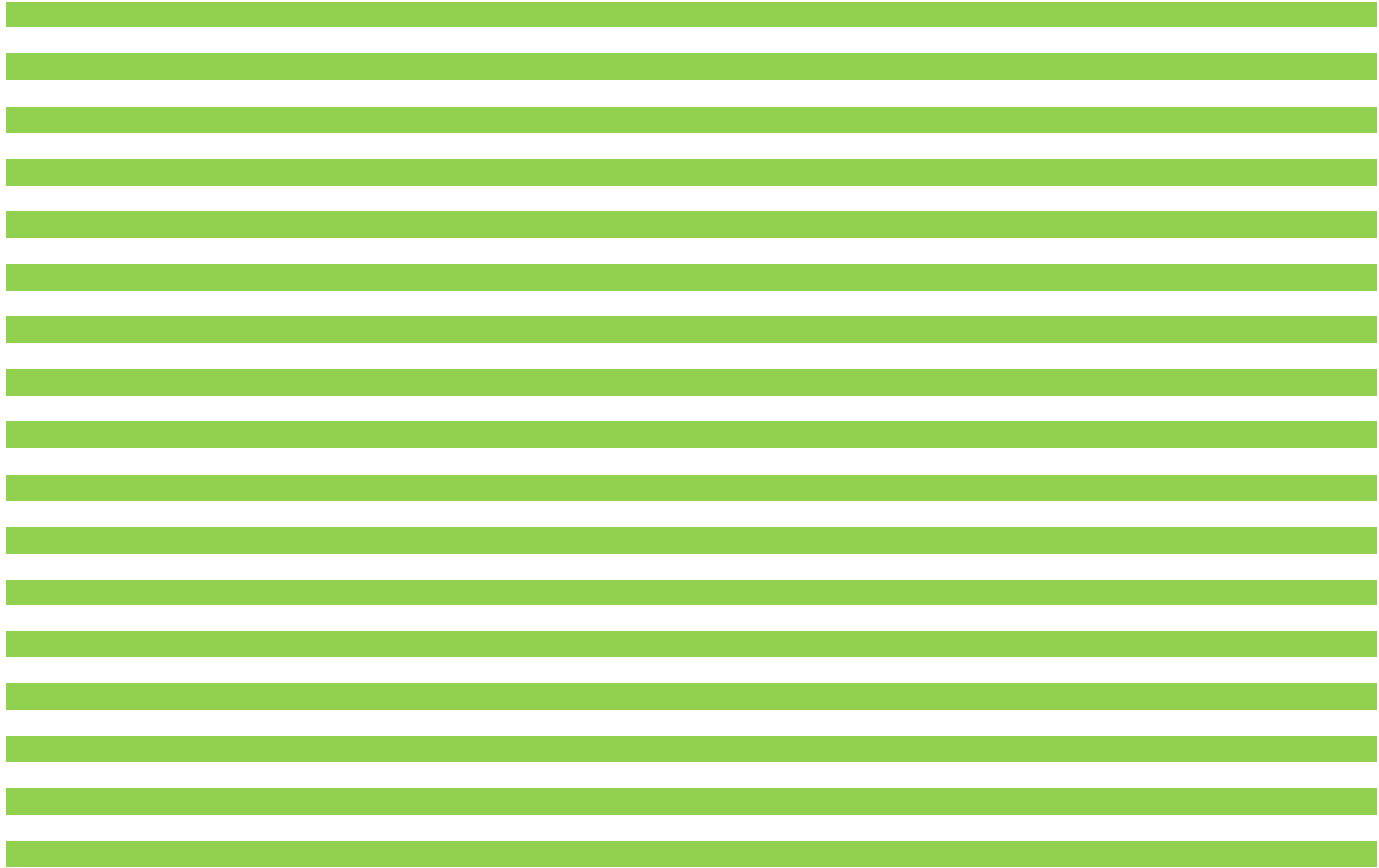
TAB]



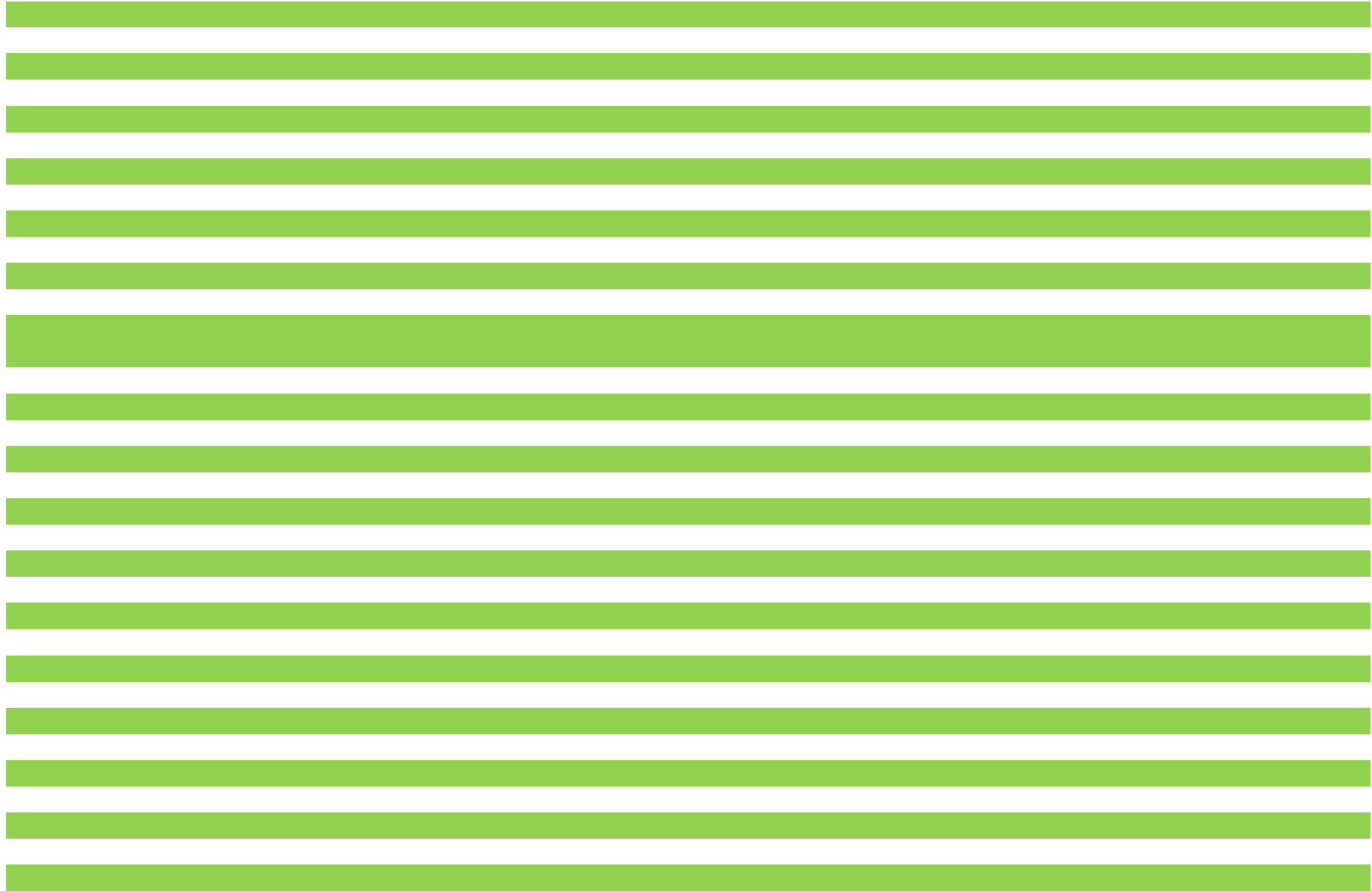
TAB]



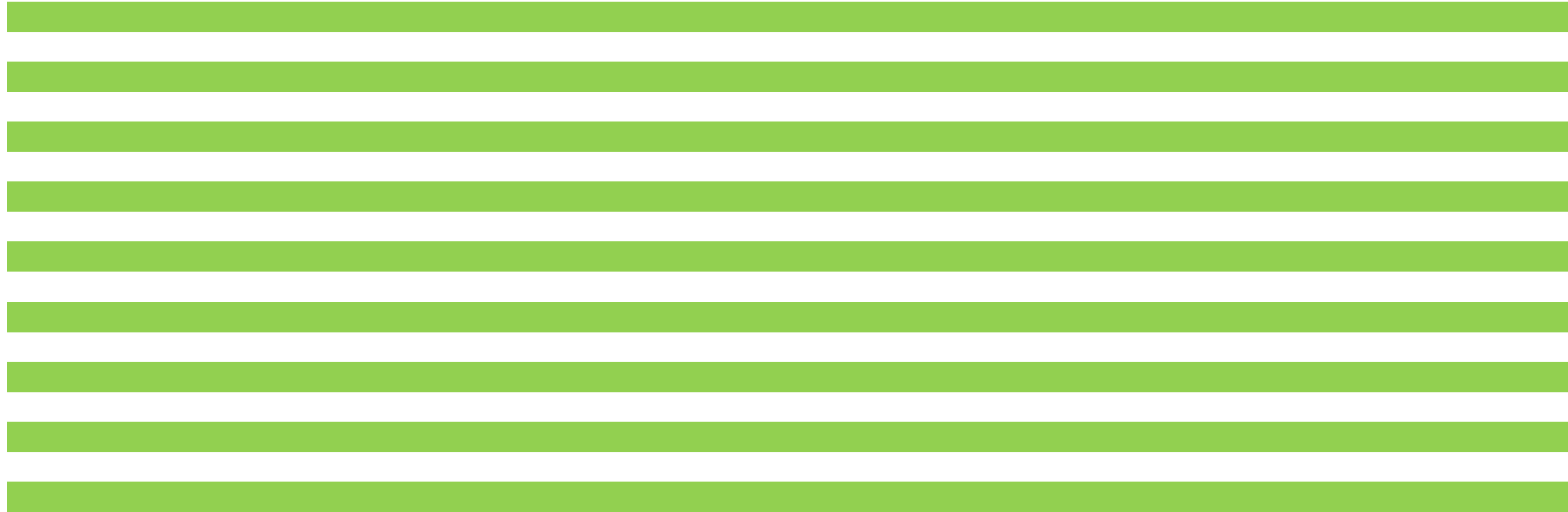
TAB]



TAB]



TAB]



TAB]

TAB]

TAB]

TAB]

TAB]

TAB]

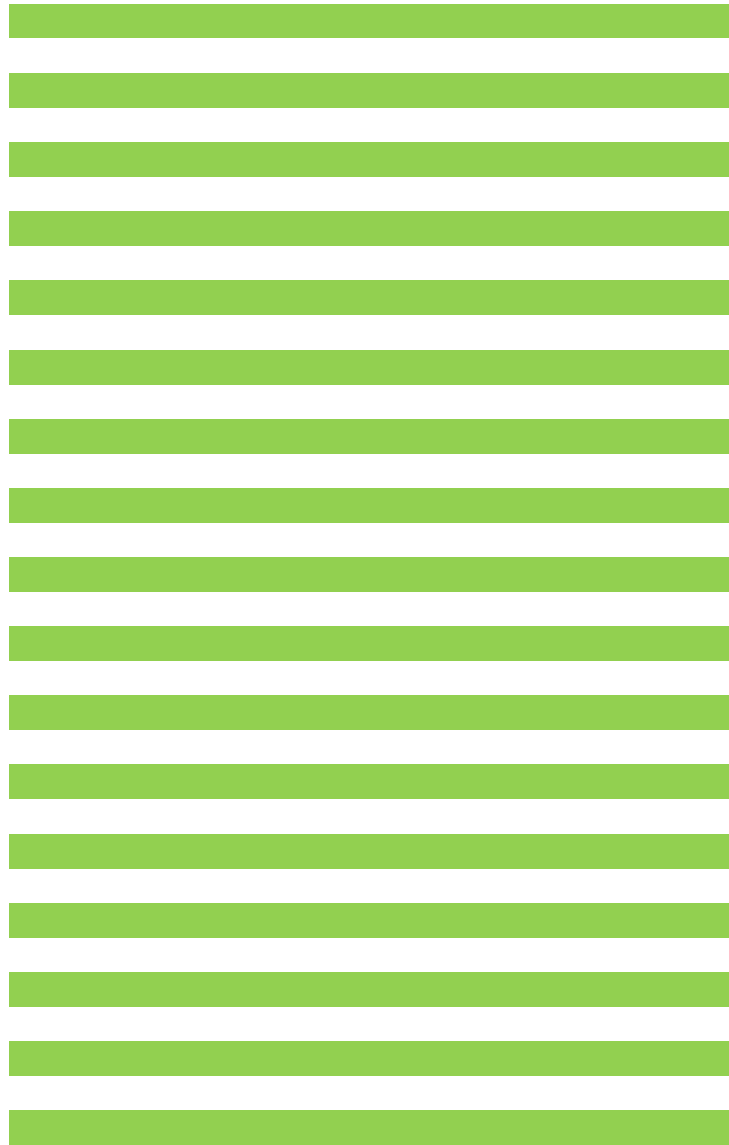
TAB]

TAB]

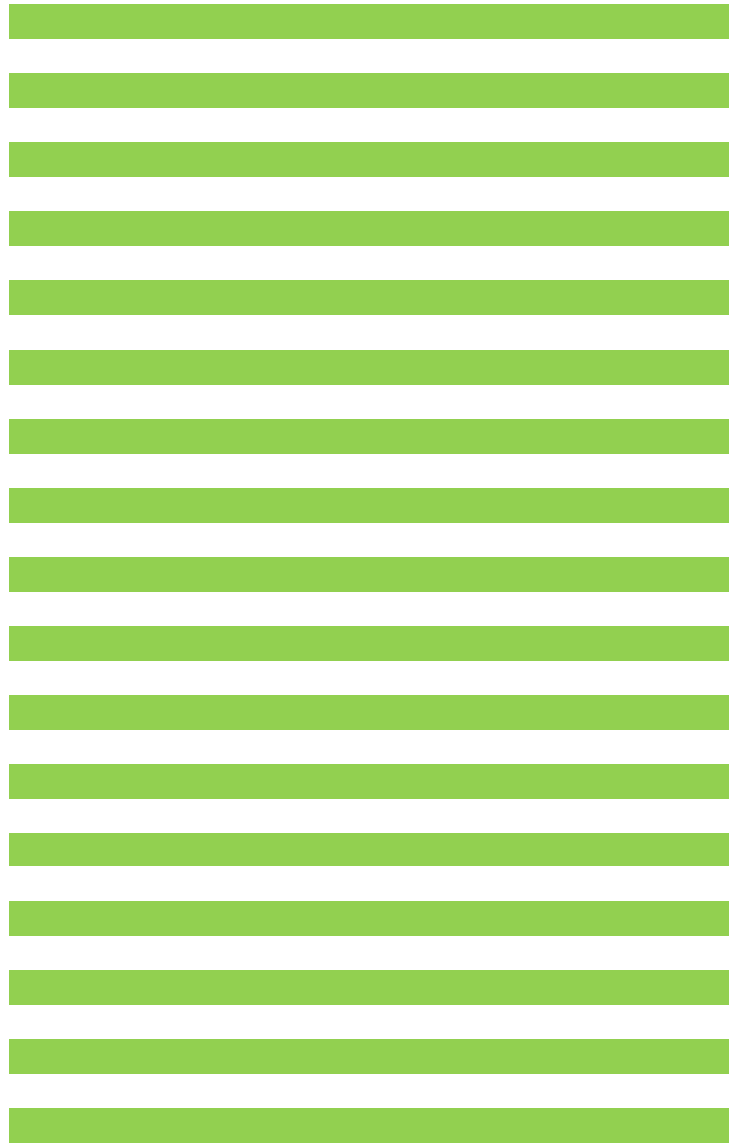
TAB]

TAB]

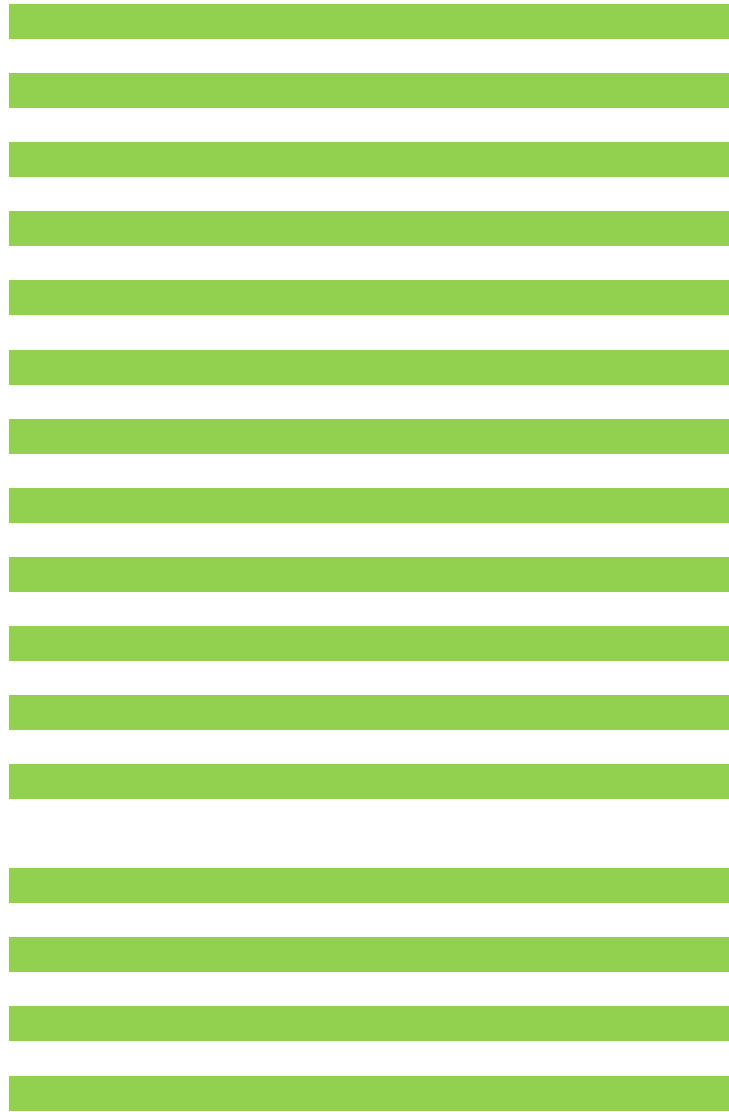
TAB]



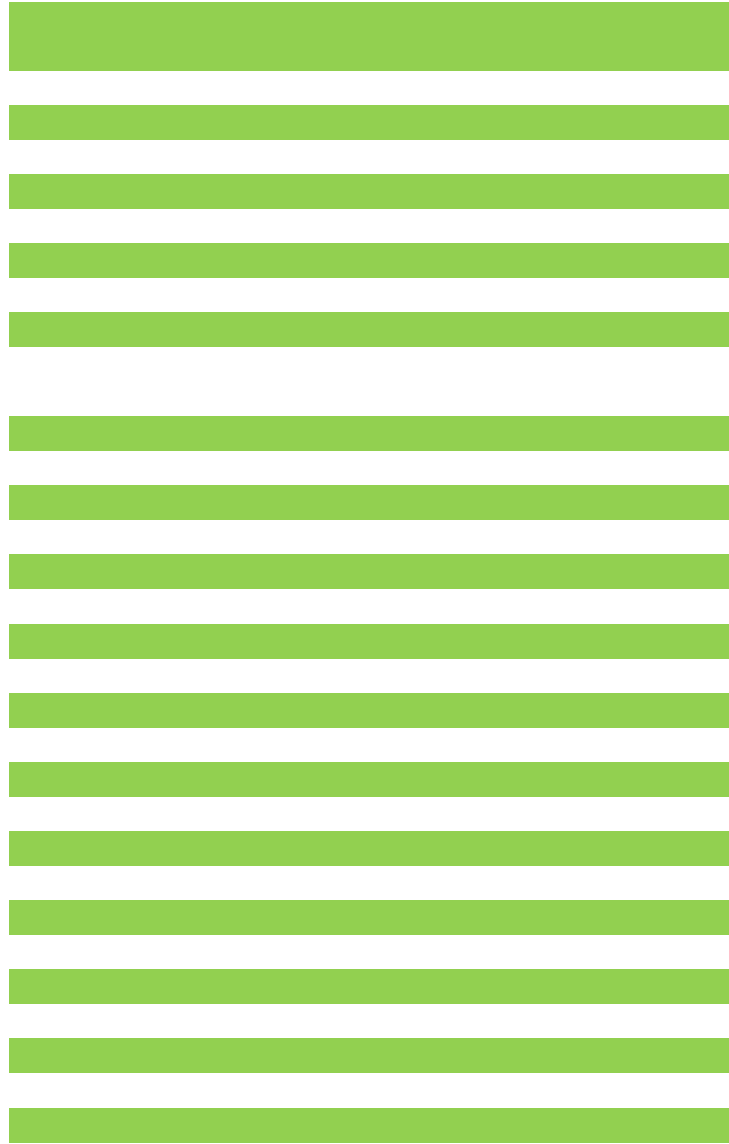
TAB]



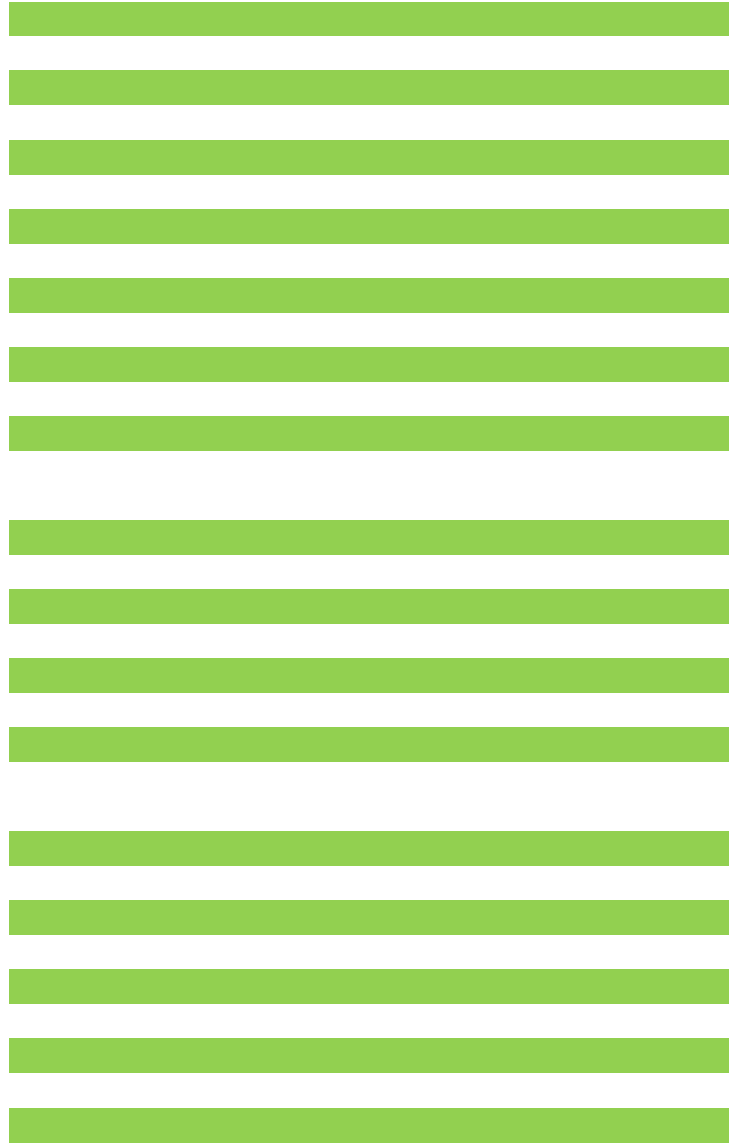
TAB]



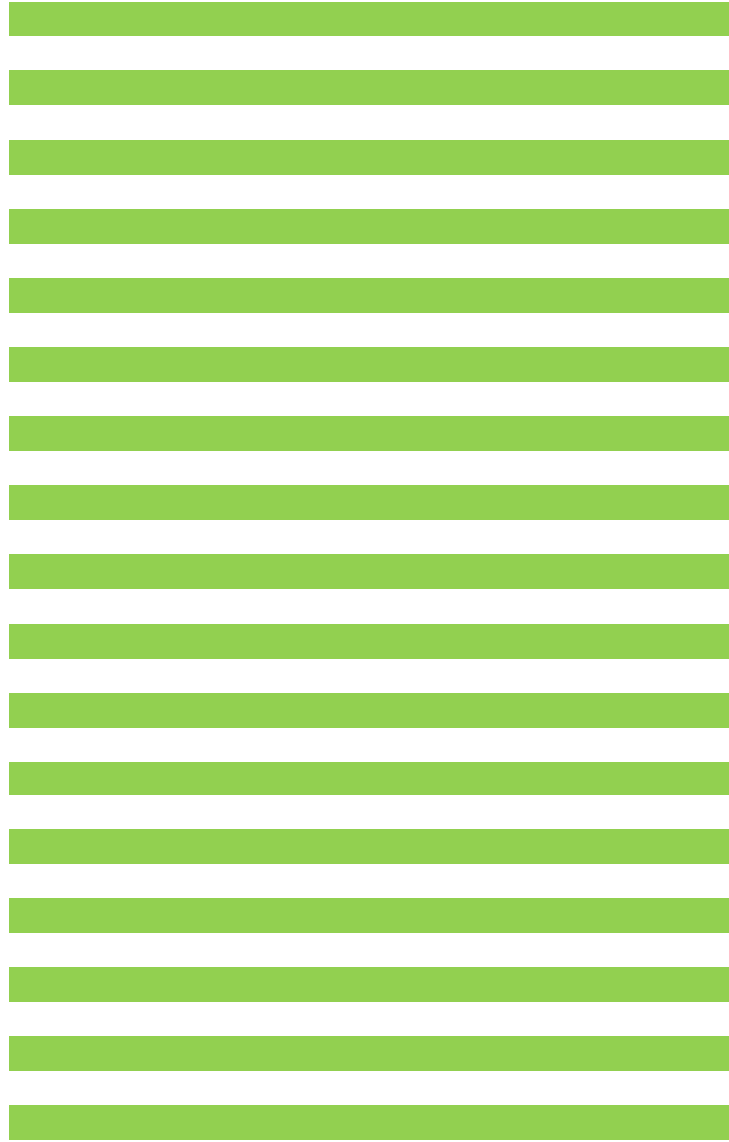
TAB]



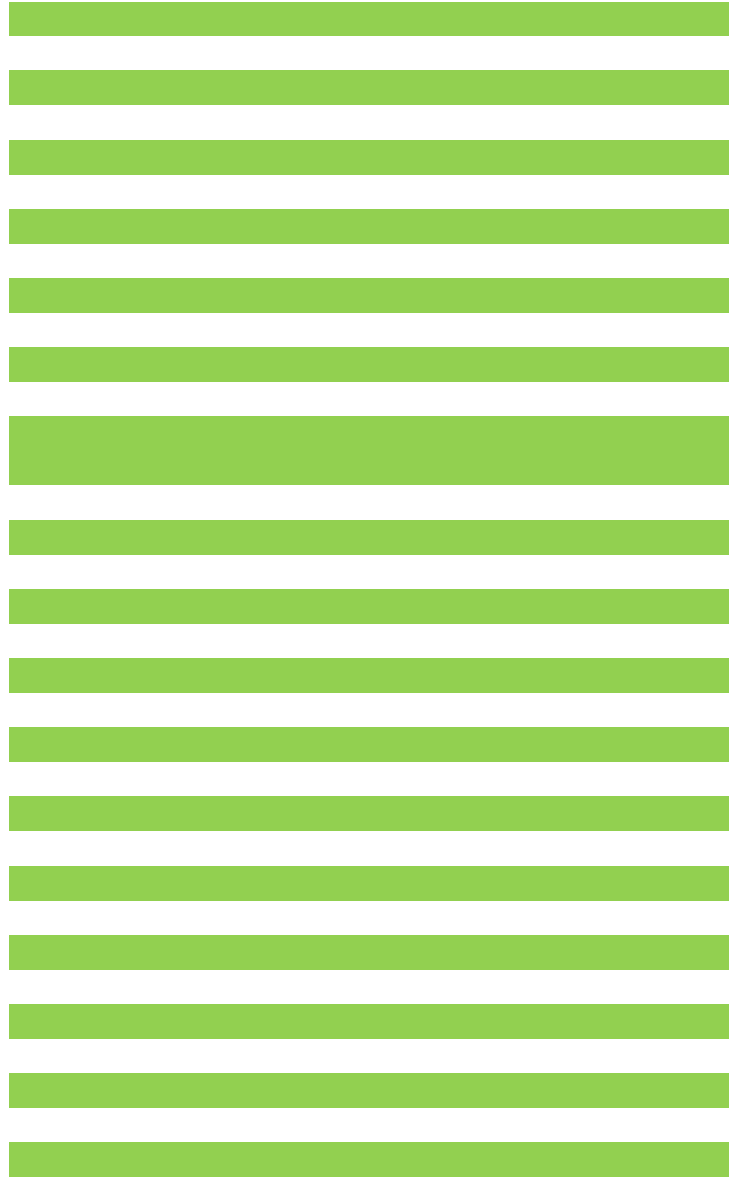
TAB]



TAB]



TAB]



TAB]

