

Results: Bloc de notas									
Archivo	Edición	Formato	Ver	Ayuda					
Gene or locus name	Number of nucleotides	Primer F	Primer R	Tm Primer F	Tm Primer R	Ta Primer F	Ta Primer R		
dnaA	1341 pb	5'- TACCTTTTATATAATCTGGACA -3'	3'- TTAATTTCTTGTGCGAATTTATC -5'	56	54	52	50		
dnaN	1137 pb	5'- TACTTTAAGTGCTAAGTTTTTC -3'	3'- GGAAGGACAGTCTTGGATAATT -5'	56	62	52	58		
rlbA	216 pb	5'- TACCGTTTAGGCTAAAGTTAAC -3'	3'- AAGTAAAGTTCAGCAGTTAATT -5'	60	56	56	52		
recF	1113 pb	5'- AACATATAGGTCCTTGAATCTTG -3'	3'- TTTACCACGCAATCACTTCACT -5'	58	62	54	58		
remB	246 pb	5'- AACATATAAGTAAATCCACTAC -3'	3'- ATACAAACTTTATCTAAGAATC -5'	56	54	52	50		
gyrB	1917 pb	5'- TACCTTGTCTGCTGTTTTGTCAA -3'	3'- GCAATTTTAGAAGCTGATAGATT -5'	62	56	58	52		
gyrA	2466 pb	5'- TACTCACTTGTTTTGTGTGGTG -3'	3'- TCTTCTTGTCTCTTCCACACT -5'	62	60	58	56		
yaaC	948 pb	5'- TACTGTATAGTACTCACCTTTC -3'	3'- TGTCAAAGAAGAGTAAAAATT -5'	60	54	56	50		
guaB	1467 pb	5'- TACACCCCTTTCATTTAAAAATT -3'	3'- TGGATTGATATGTTAAAGATT -5'	56	54	52	50		
dacA	1332 pb	5'- AACCTTGTAGTTCTTTACATTGG -3'	3'- ATGCCATTGGCCGACCAAAATT -5'	56	64	52	60		
pdxS	885 pb	5'- TACCGAGTTTGTCCATGACTTG -3'	3'- ATACGTTCTTGCGCCGACCATT -5'	64	66	60	62		
pdxT	591 pb	5'- TACAATTGTTATCCACATGATC -3'	3'- ATTCGTTTTCCGTGAACATATT -5'	58	58	54	54		
serS	1278 pb	5'- TACGAACATAGCTTTTACGACT -3'	3'- ATCTCTTCATTACTTTGGCATT -5'	60	58	56	54		
dck	654 pb	5'- TACTTCCTTGTAGTATAGGGAT -3'	3'- AGTACTATTAGTTTTTTTCACT -5'	60	54	56	50		
dgk	624 pb	5'- TACGTATGTGCGGGAATAAAC -3'	3'- ACATTTCTCGCAATAGCATACT -5'	60	60	56	56		
sleL	1284 pb	5'- CACCATTTTGTTCGCCGTGTGAG -3'	3'- ACAGTTTTTCTGCAAAAGCTATT -5'	66	58	62	54		
yaaI	546 pb	5'- AACAGGTTTCGTCTATTTCCGGG -3'	3'- TCTTGTTTAATATCTTTTTATT -5'	64	50	60	46		
tadA	486 pb	5'- TACTGTGTTCTACTTGAAATAT -3'	3'- GTCCTTTTTGAACAGACTATTC -5'	56	60	52	56		
dnaX	1692 pb	5'- CACTCAATGGTTCGAAATATAG -3'	3'- AAATTAACCTTTATTTTCTGATT -5'	60	50	56	46		
ebfC	324 pb	5'- TACGCACCGCCTTACCATTAT -3'	3'- CTTGTACGGTCCAAATAAGATC -5'	66	62	62	58		
recR	597 pb	5'- TACGTTATAGGACTTGGTTATA -3'	3'- ACTTCCTCTGCACTTAACATT -5'	58	60	54	56		
yaaL	225 pb	5'- TACCCAAAAGAAGCGTCTCTTT -3'	3'- TTTTGTAGTCGCCACCTTCATT -5'	60	62	56	58		
bofA	264 pb	5'- TACCTCGGATAAAAAATAAAC -3'	3'- GTAATTCGTTAAATAGTAAATT -5'	56	52	52	48		
csfB	195 pb	5'- TACCTGCTTTGTCAATTTGAAT -3'	3'- ATGTGGCGGAATATAAGATATC -5'	58	60	54	56		
xpaC	615 pb	5'- TACGTCTCTAAAAAGTGAAAA -3'	3'- TTTTGTAAAGTATTTCTTTATT -5'	56	50	52	46		
yaaN	1161 pb	5'- TACTTGGCTCTGTTTTCGCTGG -3'	3'- CTAATGTTATTTTCTTTTCTACT -5'	68	56	64	52		
yaaO	1443 pb	5'- TACTTGTGTGAAATATATTTC -3'	3'- ATATCTTCTCTCTTTAGTACT -5'	56	58	52	54		
tmk	639 pb	5'- TACTCGCCAAATAAATATGTA -3'	3'- CAACTTTTTTTAAAGTTAACACT -5'	56	54	52	50		
darA	330 pb	5'- TACTTTAACTATCACCCTCAAC -3'	3'- GCTACTTAAAGTAGTTAAAAATT -5'	60	54	56	50		
yaaR	441 pb	5'- CACTTTTAAATTTTCTATATG -3'	3'- TAATTAAATGGAAATGTGATC -5'	52	54	48	50		
holB	990 pb	5'- TACCGTTATAGGACCTTCTCTT -3'	3'- CAATTACACAGCTCCTCCCTATT -5'	62	62	58	58		
ricT	828 pb	5'- AACATGTTTACATTAAACCACAGG -3'	3'- TAGGCGTGTTTGGTGTCTAATT -5'	60	62	56	58		
dnaH	360 pb	5'- AACCTATTTTTTCTCAATAAAC -3'	3'- CAGTAAGAATTTATTTTTTATC -5'	54	52	50	48		
trmNF	744 pb	5'- TACCAAAGTAATGTACTACTAC -3'	3'- GTAAAAATACCTCTGTTTATT -5'	58	54	54	50		
yazA	300 pb	5'- TACCTCTGTTTATTTGGTAAAGA -3'	3'- ATTCGCGGTTTACTCGAAATT -5'	58	62	54	58		
rsmI	879 pb	5'- TACAATTCCGCGGTTTACTCGA -3'	3'- ACGAATGGTATAATTCTGTATT -5'	64	56	60	52		
abrB	291 pb	5'- TACAAATACTTTAGATGACCAT -3'	3'- GGTGGAAGTTTGGAAATTTATT -5'	56	58	52	54		
metS	1995 pb	5'- TACGGCGTTCTTTTGTATGTA -3'	3'- TTTTCCGTGTTCTTAATTTATT -5'	60	54	56	50		
dayD	768 pb	5'- TACAACAAACTGTGAGTGC GCG -3'	3'- TTCTGAAAAGGCATATTTGACT -5'	66	58	62	54		
yabE	1314 pb	5'- AACCTCTTTTCTCTTCCCTCCC -3'	3'- GTTTTAAATTTTAGAATTTAATC -5'	68	50	64	46		
rnmV	561 pb	5'- TACTTTTAAATTCTCTAGTAAC -3'	3'- CGCCCTCCTTGTCTTACTTATT -5'	54	64	50	60		
rsmA	879 pb	5'- TACTTATTTCTATAACGCTGTG -3'	3'- TAACATATTTTCGGGAAAAGATT -5'	58	56	54	52		
prtG	873 pb	5'- CACGTTAAAGTTTATCCCTCAT -3'	3'- ATACGGAATATTCAGGTTAATT -5'	60	56	56	52		
veg	261 pb	5'- TACCGCTCTCGCAACAGGCTAT -3'	3'- GCGTTCGAGTCAACGCGAAAATT -5'	66	64	62	60		
sspF	186 pb	5'- AACCCGTCGCGAGCACTCAAT -3'	3'- AGTCTTAGCTCTTATTGGCTATT -5'	68	58	64	54		
ispE	870 pb	5'- TACGCATAAAATCTTTTTCCGCG -3'	3'- GCTTGTCTTGGGAGAACTAATT -5'	60	62	56	58		
purR	858 pb	5'- TACTTCAAAGCAGCTCGCCGT -3'	3'- CTTACCTCTCTGTCTTAGTACT -5'	68	62	64	58		
ridA	378 pb	5'- TACTGTTTTCTGTCAGGTGTGTT -3'	3'- TCAATAACGTGACCATTTTATT -5'	62	58	58	54		
spoVG	294 pb	5'- CACCTTCAATGACTGCATTCTA -3'	3'- GCTTCTTCGACCTCGAAGAATT -5'	62	64	58	60		
glmU	1371 pb	5'- TACCTATTCCGCAACGCTCAAC -3'	3'- CTTTTATAAGTATTTTTTATT -5'	64	48	60	44		
prs	954 pb	5'- TACAGATTAGTTATGCTCTAT -3'	3'- TCAGTCGATAGACAAGTCGATT -5'	58	62	54	58		
ctc	615 pb	5'- TACCGTTGAAATTTGCGGTTTTTC -3'	3'- ACTTCCACTTTTGTCTGTTATT -5'	62	58	58	54		
pth	567 pb	5'- TACGAACACTAACGGCCAGAGC -3'	3'- GCTTAAATTGGGTTTCCATATT -5'	68	58	64	54		
fin	231 pb	5'- TACCGAAACGTAATAATAACAG -3'	3'- TATAGTGTGTAAATAAGTTATT -5'	58	52	54	48		
mfd	3534 pb	5'- TACCTGTTGTAAGTTTGGAAAT -3'	3'- CGTTTGGTAAAGTAGTTGCAAT -5'	58	60	54	56		
spoVT	537 pb	5'- TACTTTTCGTTGGCCATAGCATG -3'	3'- CGGAGCAGTTTACCTTGTCAATC -5'	64	66	60	62		
yabM	1599 pb	5'- TACCTGCTAAGTTATCCGCAAT -3'	3'- CTCTCTTGTTTACCGCCCAATT -5'	62	64	58	60		
yabN	1470 pb	5'- TACCGCCCACTTTTAATGTCAAG -3'	3'- TCTTTGACTCTCTCTAGTATC -5'	64	62	60	58		
hslR	261 pb	5'- TACTCTAATCTATTTAAAAACT -3'	3'- TCTTCTTTTTGAGCCGCTTATC -5'	52	62	48	58		
spcP	303 pb	5'- TACTTAAGTATAATACTAGTTT -3'	3'- AAAATCGTTCACAAATTTACT -5'	52	54	48	50		
spcQ	636 pb	5'- TACTGCGACTGCTGTGTTAAGA -3'	3'- GTGTGCCAAAACTTCTCTACT -5'	64	62	60	58		
divIC	378 pb	5'- AACTTAAAAAGGTCCCTTGCTT -3'	3'- ACACCTCTCTTCTCTGTTCAATC -5'	60	64	56	60		
yabR	387 pb	5'- TACAGCTAACTTCAACCGTCGT -3'	3'- CGCGCCTCGTCTCTCTCTATT -5'	64	68	60	64		
spoIIE	2484 pb	5'- TACCTTTTTCTGCTCTTTCTCTC -3'	3'- TTTGTTTGTCTTTAAAGTATT -5'	60	52	56	48		
yabS	738 pb	5'- CACTTGTGCTCTGTAATTTTAG -3'	3'- GAAACTGCTCAACCCGTACATC -5'	60	64	56	60		
prkT	1017 pb	5'- TACTACTTGTGCGAAACTGCT -3'	3'- ATAATAGAAAAAGAAATAGACT -5'	64	52	60	48		
tilS	1419 pb	5'- CACTTTTACAGTTTCTAAAAA -3'	3'- ATCCCCGTTCTGTTTTAGTACT -5'	56	64	52	60		
hprT	543 pb	5'- TACTACTTTGTACTATAAGCTCT -3'	3'- TGGCCGTCAAACTTTTCGACT -5'	58	64	54	60		
ftsH	1914 pb	5'- TACTTAGCCCAAGAGGCATTAT -3'	3'- CTTTCTGCTATGCTTTCTCATT -5'	62	60	58	56		
coaX	777 pb	5'- AACCAATGACCAATAGCTACACC -3'	3'- TTTGGCGCATCTTCCACATATC -5'	62	64	58	60		
hslO	876 pb	5'- TACCTAATAAATCATTTTCGTG -3'	3'- AGCACTGGTTTGATGGGCGATT -5'	56	66	52	62		
yacD	894 pb	5'- AACTTTAGTCTCTGTTAGACCT -3'	3'- ACCCTTTTTCATGATTCACT -5'	58	60	54	56		
cysK	927 pb	5'- TACCATGCACTCGTTTGAGGT -3'	3'- CGGCGACATGGTTAAGCTAATT -5'	64	64	60	60		
pabB	1413 pb	5'- TACCGTGTTCGCTCTGGCCGTC -3'	3'- TCTTCTCTGTTTTAATCTACT -5'	72	56	68	52		
pabA	585 pb	5'- TACTAAAAATACTAACTATTGA -3'	3'- AGCGTTCCTCAATAACGCACCT -5'	52	64	48	60		
pabC	882 pb	5'- TACTATATACACTTGC CGGCCA -3'	3'- TCGTACCGCGCTGTGTGTTATC -5'	64	68	60	64		
folP	858 pb	5'- TACCGCGTCGTGTGTTATCTAG -3'	3'- CCTCCCCACGTGGTACCTATT -5'	66	70	62	66		
folB	363 pb	5'- TACCTATTTCAAATACATCTTC -3'	3'- TTAATGCTCTTTTTTTCAGTACT -5'	56	56	52	52		
folK	504 pb	5'- TACTTGTGTATCGAATAATAAC -3'	3'- TAAGCACGTAAAGTCTTTCGACT -5'	56	62	52	58		
yazB	210 pb	5'- TACCTTCGTTTATAGACCCCAT -3'	3'- TAACCGCGCGGTCTTTTTATC -5'	66	66	60	62		
dusB	1002 pb	5'- AACAAAGTTTTAGCCTCTATAAG -3'	3'- AGTTTACGTTTTTATCTCTATT -5'	58	56	54	52		
lysS	1500 pb	5'- TACTCAGTTCTTCTCGTATTGG -3'	3'- GGGCGTTTACTCTGTGCGATT -5'	62	66	58	62		
ctsR	465 pb	5'- CACCCTGTATTATAAAGACTGT -3'	3'- CTGAAGAAATTTAAATTTTATT -5'	60	50	56	46		
mcsA	558 pb	5'- AACTAAACAGTTCTCACGGTGC -3'	3'- ATCACTCTCCTTGTCTCTATT -5'	64	64	60	60		