

Table S4. Diversity in the ampliconic and low diversity regions. The table shows the pi value computed for the nonPAR region of the X chromosome. This compares the nonPAR, nonPAR with ampliconic regions filtered out, nonPAR with low diversity regions filtered out, and nonPAR with both sets of regions filtered out. It also shows the pi values for each ampliconic and each low diversity region.

Region	Chr X position	Pi
nonPAR	2700000-154940559	0.00051179
Ampliconic region 1	48202745-48292983	0.00000447
Ampliconic region 2	48976199-49062381	0.00000678
Ampliconic region 3	51395467-51492862	0.00000235
Ampliconic region 4	51775560-51966529	0.00000402
Ampliconic region 5	52518132-53027386	0.00001449
Ampliconic region 6	55464117-55574172	0.00000691
Ampliconic region 7	62335733-62495350	0.00000067
Ampliconic region 8	70894117-71055682	0.00000457
Ampliconic region 9	71941159-72325075	0.00001086
Ampliconic region 10	100818723-100903977	0.00003235
Ampliconic region 11	101435778-101774391	0.00002782
Ampliconic region 12	103195105-103362341	0.00013457
nonPAR- ampliconic regions	--	0.00051647
Low diversity region 1	10241177-12619185	0.00009100
Low diversity region 2	16946047-18747389	0.00005870
Low diversity region 3	19303480-22198160	0.00008547
Low diversity region 4	38344992-41272675	0.00002217
Low diversity region 5	45930478-77954462	0.00008178
Low diversity region 6	99459295-111145964	0.00007904
Low diversity region 7	128232540-136796526	0.00011957
Low diversity region 8	151519514-155156362	0.00019179
nonPAR- low diversity regions	--	0.00060150
nonPAR-ampliconic and low diversity regions	--	0.00060150