

Fusion/Gene	Left Primer	Left Sequence	Right Primer	Right Sequence	Size (bp)
A	4L-29	CAGGAAGAGCTGCTGGAAAC	4R-29	AGTGAATAGGCCAGGTGTGG	181
B	2L-29	GCTTGGGTGATCTGGAAAAG	2R-29	AGATGGCCACTTGGAGACAC	202
C	1L-29	CAGCTATTTGGGAGGCTGAG	1R-29	GGCAGTGGAAATCTGTGTGTG	762
D	8L-29	CTGGGGGAACCTTCAATTTT	8R-29	AAAAAGGCACAGCTGGAGAA	193
E	5L-29	CAGGCACTTGCCATTCATA	5R-29	TCCAGTGGCAACTACCCTTC	179
F	3L-29	TGAGGCCCTGTCTCAAATAAA	3R-29	ACACACGTGGATGCAGACAG	170
G	9L-29	GGGGAATATTTGCAGTGTGG	9R-29	GAAATTCAGCCCCCTCAGTT	206
H	6L-29	TCTATGCCAGTGCTTACTCCTT	6R-29	CTGTTCCACCAAAGCCAAAT	155
I	7L-29	TGAGCAGATTTTCTGTATATTTT CA	7R-29	GTCTCCCAGGCTGCAGTG	150
GPX7	GPX7 Left	ATTCTGGTTGGAGTGGTCTG	GPX7 Right	AAAAGGAAGGAGAGCAAAGC	129