

Table S1
Overview probe sequences of the KIR MLPA technique.

Probe	Sequence (5'-3')
2DL1 - #1 left	GGGTTCCCTAAGGGTTGGACTACCCCATCGCTCTTCATGCTGGATCATTCACTCTGCATCCCAATGACAATG
2DL1 - #1 middle	AGAAGAAAGTCTGGACACTCTCACCTATGATCACGATGTCAGAGGGTCACTGGGAGCTGACAC
2DL1 - #1 right	CTGATAGGGGGAGTGAGTAACAGAACCGTAGTCTAGATTGGATCTTGCTGGCAC
2DL1 - #2 left	GGGTTCCCTAAGGGTTGGACATCCTGTGCGCTGCTGAGCTGAGCTCG
2DL1 - #2 right	GTGCGGGCTGCCTGTCTGCTCCGGCAGTCTAGATTGGATCTTGCTGGCAC
2DL2 - #1 left	GGGTTCCCTAAGGGTTGGACTGACCTTGGGCCCTGCAGAGAACCCTACA
2DL2 - #1 right	TTCATGGGCCTCCCCCTCCCTGGATGTCTAGATTGGATCTTGCTGGCAC
2DL2 - #2 left	GGGTTCCCTAAGGGTTGGACATGTCTATGATCCTAGAGCCTTAGCTGAGGAGCTTCTGCTGATGATGGAGAT
2DL2 - #2 right	AAGCATGGACAGATGCAGAGAGAAGACGAAGCTTGGGTGTGAGGGAGGTCTAGATTGGATCTTGCTGGCAC
2DL3 - #1 left	GGGTTCCCTAAGGGTTGGACYRCACAGTTGRATCACTGCGTTTTACACAGAGAAAAATCACTCRCCCTT
2DL3 - #1 middle	CTCAGAGGCCAAGACACCCCCAACAGATATCATCGTGTACACGGAACTTCCAAATGCTGAGCCCT
2DL3 - #1 right	GATCCAAAGTTGTCTCCTGCCATGAGCACCACAGTCAGGCCTTGTCTAGATTGGATCTTGCTGGCAC
2DL3 - #2 left	GGGTTCCCTAAGGGTTGGACTGCTGCCTTGGGCCAGGGACCATCCTGTCTGTGAGGAACACACACCTGAGTGCTC
2DL3 - #2 middle	CCATCCTGCTTCCCACATGGCCCTGAGCTCTCTGGCCTCTGCTTCGTGAGACTTACTTTTTTTGTTGC
2DL3 - #2 right	AGCACCAGCGATGAAGGAGAAAGAAGAGGAGGAGGATGAAGAGGATGTCTAGATTGGATCTTGCTGGCAC
2DL4 - #1 left	GGGTTCCCTAAGGGTTGGACCTGCTTCAGAACATGGCTCTCTGCTGGGGAGACACCCAA
2DL4 - #1 middle	TCTGCAGGCCATAGTGAACCCTGGTGTCTCCTTCCCTTCCAGGACTCACCAAG
2DL4 - #1 right	ACATGCCAGGATGATGACCGTGGGTGACATGGAGTCTAGATTGGATCTTGCTGGCAC
2DL4 - #2 left	GGGTTCCCTAAGGGTTGGACCCGCTTAGAAAGAAGAAATGGGGAGAATCTTCTGAGCACAGGGAGGGAGGGGC
2DL4 - #2 middle	AGCTCAACATACTCCTCTCTGAGGCGGCATCTCCTTCTCCCAAGGTGGTCAGGACAAGCCCTTCTGC
2DL4 - #2 right	TCTGCCTGGCCCAGCGCTGTGGTGCCTCAAGGAGGACACGTGACTTTCGGTGGTCTAGATTGGATCTTGCTGGCAC
2DL5 - #1 left	GGGTTCCCTAAGGGTTGGACTCAGGTGTGAGGGGAGCTGTGACAAGGAAGAACCCTCC
2DL5 - #1 middle	CTGAGGAAACTGCCTCTTCTTCCAGGTCTATT
2DL5 - #1 right	TGGGAAACCTTCACTCTCAGCCCAGCCGGTCTAGATTGGATCTTGCTGGCGC
2DL5 - #2 left	GGGTTCCCTAAGGGTTGGACCGACCTACACATGCTTYRGCTCTCTCCATGACTC
2DL5 - #2 right	ACCCTATGAGTGGTCAGACCCGAGTGACCCGTCTAGATTGGATCTTGCTGGCAC
2DL5b left	GGGTTCCCTAAGGGTTGGACATGTTAGCACAGATTTTAGGCATCTCGTGTTCGGATAAAAATACATGAAAAGTCTTTCAC
2DL5b middle	GTTAGCACAGATTTTAGGCATCTTGTTTCGGGAGGTTGGATCTGAGACGTGTTGTGAGTTGGTCATAGTGAAGGACGT
2DL5b right	GAGGTGCCAATTCTAGTGAACAATTTCCAGGAAGCCGTGTTCCGGTCTAGATTGGATCTTGCTGGCAC
2DP1 - #1 left	GGGTTCCCTAAGGGTTGGACCCAAGGTGGTCAGGACAAGCCCTTCTGCTGCCTGGCCCAGCTC
2DP1 - #1 right	TGTGGTGCCTCCAGGACATGTGATTCTTCCGGTGTCTAGATTGGATCTTGCTGGCGC
2DP1 - #2 left	GGGTTCCCTAAGGGTTGGACCAGGGACCTACAGATGCTACGGTCTGTTACTCACTCCCC
2DP1 - #2 right	CATCAGTTGTAGCTCCCAGTGACCCTCTGGACATCGTCTAGATTGGATCTTGCTGGCAC
2DS1 - #1 left	GGGTTCCCTAAGGGTTGGACACAGGGCCCATGAAAAGGCTGTTCCAGAATATTATGTTGTAGAGCTCAGGGACAGGCA
2DS1 - #1 middle	CCCCATCTTCTTTACAGACTGAAGTTGTTAAACCAAGATAAGAATGACACTGAAGAATCACATA
2DS1 - #1 right	TCCTGGAGGCACCACAGGGCTTGGCCAGTCTAGATTGGATCTTGCTGGCAC
2DS1 - #2 left	GGGTTCCCTAAGGGTTGGACTAGGAGACCGTGGAAAAGGCAATCCCGA
2DS1 - #2 middle	CCCCTGGTGAATGTGGTGTCTGATTTT
2DS1 - #2 right	GACACTAAGTGGATGAAGCAGATGGATATAAGCGTCTAGATTGGATCTTGCTGGCAC
2DS2 left	GGGTTCCCTAAGGGTTGGACCCGGCCGAGCACCCAGGGTCTCTTCCC
2DS2 right	AGTTTATGAGAGACTCCCTGACAGGACGTCTAGATTGGATCTTGCTGGCAC
2DS3 - #1 left	GGGTTCCCTACGGGTTGGACCATCACGATGTCCAGAGGGTCACTGGGAGCTGAA
2DS3 - #1 right	AACTGATAGGGGGAGTGAGGAACAGAGACCGTCTAGATTGGATCTTGCTGGCAC
2DS3 - #2 left	GGGTTCCCTAAGGGTTGGACCGAAAGACCCGAAGCATCTGTAGTTCTCTCT
2DS3 - #2 right	TGGGTGGCAGGGCCAGAGAAAGTCGGCCTGGAATGTCTAGATTGGATCTTGCTGGCAC

Table S1 continued
Overview probe sequences of the KIR MLPA technique.

Probe	Sequence (5'-3')
2DS4 - All left	GGGTTCCCTAAGGGTTGGACTCCCCTCTCTGTGCAGAAGGAAGTCTCAAACATGACATCC
2DS4 - All right	GACCAACATTGCAGGATGACTGTCTCTTCTGATTTACCAGGTGACCTGGGAGTCTAGATTGGATCTTGCTGGCAC
2DS4 - WT/trunc left	GGGTTCCCTAAGGGTTGGACTTGGGCCAGAGGAAAGTCRGCCTGGAATGTTCCGKGTAT
2DS4 - trunc middle	GCTGCGCACTGCAGGGAGCCTACGTTTCATGGGCCTCCCCYTCCCTGGATAG
2DS4 - trunc right	ATGGAGCTGCAGGACAAGGTACAGTCTAGATTGGATCTTGCTGGCAC
2DS4 - WT middle	GCTGCGCACTGCAGGGAGCCTACGTTTCATGGGCCTCCCCYTCCCTGGATAGATGGTAC
2DS4 - WT right	CATGTCATAGGAGCTCCGGGAGCTGCAGGACAAGWCACATTCTCTCTAGATTGGATCTTGCTGGCAC
2DS5 left	GGGTTCCCTACGGGTTGGACCGAGTAAACCGGAAAATTTTCATCTGCACAGAGAGGGGACGTTAAC
2DS5 middle	CACACTTTGCCCTCATTGGAGAGCACATTGATGGGGTCTCCAAGGG
2DS5 right	CAACTTCTCCCTCGGTGCGATGACACAAGACCTGGCATAGCGAATACGTCTAGATTGGATCTTGCTGGCAC
3DL1 - #1 left	GGGTTCCCTAAGGGTTGGACCCCTCAHGCCTCGYTGGACA
3DL1 - #1 middle	GATCCATGATGGGGTCTCCAAGGCCAATTTCTCCATCGGTCCCATGATGCT
3DL1 - #2 left	GGGTTCCCTAAGGGTTGGACTCAGCTCAGGTATGAGGGGAGCTATGACAAGGAAGAACCT
3DL1 - #2 middle	CCCTGAGGAAACTGCCTCTTCTCCTTCCAGGTCC
3DL1 - #2 right	ATATGAGAAACCTTCTCTCAGCCAGCCGGGTCTAGATTGGATCTTGCTGGCAC
3DL1/S1 right	TGCCCTTGACGGGACCTACAGATGCTACGGTCTGGTCTAGATTGGATCTTGCTGGCAC
3DL2 - #1 left	GGGTTCCCTAAGGGTTGGAATCCACCCTAAGGTTTGGGGAKGGACTCACCCATGA
3DL2 - #1 right	GTGGCCAGGCCCCCTGCAGCAAGAAGAACCCTGTCTAGATTGGATCTTGCTGGCGC
3DL2 - #2 left	GGGTTCCCTAAGGGTTGGACCATGAAGCTCCTCAGCTATGGCTCTAGGATCATAAGACATGGGACAGACA
3DL2 - #2 middle	CGGGTTTTCTCACCTGTGACAGAAACAAGCAGTGGGTCACCTTGAGTTTGACCACACGCA
3DL2 - #2 right	GGGCAGGGCACGAAAGAGCCGAAGCATCTGTAGTTCCCTCGTCTAGATTGGATCTTGCTGGCAC
3DL3 - #1 left	GGGTTCCCTACGGGTTGGATAGATGCTTCGGCTCTTCCGTGCCCTGCC
3DL3 - #1 right	CAYGCGTGGTCAGACCCGAGTGACCCGTCTAGATTGGATCTTGCTGGCAC
3DL3 - #2 left	GGGTTCCCTAAGGGTTGGACAAGGGTGAGAGGCAGGTCTGTATTCTCTCACCTA
3DL3 - #2 middle	CGACCACGATGTCCAGAGGGTCACTGGGAGCYGACAACCTCATAGGGTA
3DL3 - #2 right	AGTGAGTGACAGAACCAAAGCATCTGTAGTCTAGATTGGATCTTGCTGGCAC
3DP1 - #1 left	GGGTTCCCTAAGGGTTGGACACCATGATCACCAGGGGGTCTGGGTGCTGACCACCCAGTGAGGA
3DP1 - #1 right	AGTGTGGGTGTGAACCCGACATCTGTAGTCCCTGTCTAGATTGGATCTTGCTGGCGC
3DP1 - #2 left	GGGTTCCCTAAGGGTTGGACGCTGAGGCCTGAAAGGAATAGAGGGAGGGAGTGCCACATC
3DP1 - #2 right	CTCCTCTCTAAGGTGGCGCCTCTTCTCCCCAGGTGTCTAGATTGGATCTTGCTGGCAC
3DS1 - #1 left	GGGTTCCCTAAGGGTTGGACTTGTTCATCAGAATCCTGGAGAGAGGAAATGCTGAGTGAGGGAGGGTGCACATTTTTC
3DS1 - #1 middle	AGGACTCTTTGGGAATAACACTAGCCACGAGGCTGGGCCAGGAGCACCTACCTCGCTGTTTAC
3DS1 - #1 right	TTCTGTTCCCTGCAGGCTCTTGGTCCATTACAGCAGCATCTGTAGGAGACGTCTAGATTGGATCTTGCTGGCGC
3DS1 - #2 left	GGGTTCCCTAAGGGTTGGACACTTCTTCTGCACAAAGAGTGGATCTCTAAGG
3DS1 - #2 middle	ACCCCTCAGCCTCGTTGGACAGATCCATGATGGGGTCTCCAAGGCCAATTTCTCCATCGGTCCATGATGCG