

Table S1. Estimated allele-level KIR haplotypes and their frequencies

KIR haplotype #	<i>2DL4</i>	<i>3DL1/S1</i>	<i>2DS4</i>	<i>3DL2</i>	Frequency
1	*001	*015	*001	*002	35.5%
2	*005/*011	S*013	Blank	*007	11.2%
3	*006	*007	*004	*008	7.2%
4	*006	*007	*004-variant	*008	5.9%
5	*001	*015	*014	*002	3.9%
6	*001	*020	*001	*009	3.3%
7	*008	*001	*003	*010	2.6%
8	*005/*011	*005	*003	*010	2.6%
9	*005/*011	*005	*007/*010	*010	2.6%
10	*001	*015	*001	*010	2.6%
11	*005/*011	S*013	Blank	*002	2.0%
12	*008	*001	*003	*001	2.0%
13	*005/*011	S*085	Blank	*007	2.0%
14	*005/*011	*005	*007/*010	*002	1.3%
15	*005/*011	*005	*007/*010	*009	1.3%
16	*006	*007	*004	*016	1.3%
17	*001	*015	*015	*002	1.3%
18	*005/*011	S*013	Blank	*015	1.3%