Identification of minor histocompatibility antigens based on the 1000 Genomes Project

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<th>Frequency (%)</th>
<th>SD</th>
<th>Minimal Frequency</th>
<th>Maximal Frequency</th>
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<td>98.72</td>
<td>2.01</td>
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<td>96.98</td>
</tr>
</tbody>
</table>

**Table 51: mtHag phenotype frequency in percentage (%) per population**

mtHags are ordered from top to bottom as they are represented from left to right in Figure 1A. The variation num. Basepair mtHag allele (Fw strand) is shown in parentheses. Immunogeneic amino acid Chromosome Position # individuals: 98 88 97 61 60 66 55 97 89 100 85 93 89 14 Average SD Minimal frequency Maximal frequency

**Statistics**

Average SD Minimal frequency Maximal frequency