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(71) Applicant: **NUCLEIX LTD.** [IL/IL]; 3 Pakeris St.,
7670203 Rehovot (IL).

(72) Inventors: **FRUMKIN, Danny**; 1 Carmel Street, 7630501
Rehovot (IL). **SAVIN, Orna**; 25 Ha Histadrut, 4642041
Herzliya (IL). **DANAN, Shacade**; 46 Ha Hagana St.,
7621515 Rehovot (IL).

(74) Agent: **WEBB, Cynthia** et al.; WEBB & CO., P.O. Box
4177, 7414003 Ness Ziona (IL).

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(57) Abstract: Methods and kits for determining the efficiency of plasma separation from whole blood are provided, which use PCR amplification or high-throughput sequencing data of genomic loci that are present at different concentrations in plasma DNA and in whole blood DNA.



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