

\*陳彥瑜博士

所有發表期刊論文

1. Chen LY\*, Wei KC\*, Huang ACY\*, Wang K, Huang CY, Yi D, Tang CY, Galas, DJ, Hood LE. RNASEQR - A streamlined and accurate RNA-seq sequence analysis program. (2012) Nucl. Acids Res. 40(6): e42 \*Contributed equally (Corresponding author)
2. International Warfarin Pharmacogenetics Consortium. Warfarin pharmacogenetics: a single VKORC1 polymorphism is predictive of dose across 3 racial groups. (2010) Blood. 115(18):3827-34
3. International Warfarin Pharmacogenetics Consortium. Estimation of the warfarin dose with clinical and pharmacogenetic data. (2009) N Engl J Med. 360(8):753-64
4. Wadelius M\*, Chen LY\*, Lindh J\*, Eriksson N\*, Ghori J, Bumpstead S, Holm L, McGinnis R, Rane A, Deloukas P. The largest prospective warfarin-treated cohort supports genetic forecasting. (2009) Blood. 113(4):784-92. \*Contributed equally ("Must read" by the Faculty of 1000 Medicine)
5. Wadelius M, Chen LY, Eriksson N, Bumpstead S, Ghori J, Wadelius C, Bentley D, McGinnis R, Deloukas P. (2007) Association of warfarin dose with genes involved in its action and metabolism. Hum Genet. 121:23-34
6. Chen LY, Eriksson N, Gwilliam R, Bentley D, Deloukas P, Wadelius M. (2005) Gamma-glutamyl carboxylase (GGCX) microsatellite and warfarin dosing. Blood. 106:3673-3674
7. Wadelius M, Chen LY, Downes K, Ghori J, Hunt S, Eriksson N, Wallerman O, Melhus H, Wadelius C, Bentley D, Deloukas P. (2005) Common VKORC1 and GGCX polymorphisms associated with warfarin dose. The Pharmacogenomics J. 5:262-270
8. Chen LYY, Lu SH, Shih ESC, Hwang MJ. (2002) Single nucleotide polymorphism mapping using genome-wide unique sequences. Genome Res. 12:1106-1111