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Research Interests: (Key words)	Breast cancer, translational research, next-generation sequencing, biomarker development, prognosis and prediction			
Selected Publications (in recent 5 yr):	 Huang CC, Tsai YF, Liu CY, Chao TC, Lien PJ, Lin YS, Feng CJ, Chiu JH, Hsu CY, Tseng LM. Comprehensive molecular profiling of Taiwanese breast cancers revealed potential therapeutic targets: prevalence of actionable mutations among 380 targeted sequencing analyses. BMC Cancer. 2021 Feb 25; 21(1): 199. Huang CS, Liu CY, Lu TP, Huang CJ, Chiu JH, Tseng LM, <u>Huang CC</u>. Targeted sequencing of Taiwanese breast cancer with risk stratification by the concurrent genes signature: a feasibility study. J Pers Med. 2021 Jun 28; 11(7):613. 			
	 Chen YJ, Huang CS, Phan NN, Lu TP, Liu CY, Huang CJ, Chiu JH, Tseng LM, <u>Huang CC</u>. Molecular subtyping of breast cancer intrinsic taxonomy with oligonucleotide microarray and NanoString nCounter. Biosci Rep. 2021 Aug 27; 41(8):BSR20211428. 			
	 Phan NN, <u>Huang CC</u>, Tseng LM, Chuang EY. Predicting breast cancer gene expression signature by applying deep convolutional neural networks from unannotated pathological images. Front Oncol. 2021 Dec 1; 11:769447. 			
	 Huang CS, Lu TP, Liu CY, Huang CJ, Chiu JH, Chen YJ, Tseng LM, <u>Huang CC</u>. Residual risk stratification of Taiwanese breast cancers following curative therapies with the extended concurrent genes signature. Breast Cancer Res Treat. 2021 Apr; 186(2):475-485. 			
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