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重要經歷:	台灣大學公衛學院流行病學與預防研究所兼任教授 台灣乳房醫學會秘書長			
研究方向: (關鍵詞)	乳癌 轉譯醫學 次世代定序 生物標記開發 預後與預測			
五年內 代表著作:	<ol style="list-style-type: none"> <li>1. <b>Huang CC</b>, 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.</li> <li>2. Huang CS, Liu CY, Lu TP, Huang CJ, Chiu JH, Tseng LM, <b>Huang CC</b>. 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.</li> <li>3. Chen YJ, Huang CS, Phan NN, Lu TP, Liu CY, Huang CJ, Chiu JH, Tseng LM, <b>Huang CC</b>. Molecular subtyping of breast cancer intrinsic taxonomy with oligonucleotide microarray and NanoString nCounter. Biosci Rep. 2021 Aug 27; 41(8):BSR20211428.</li> <li>4. Phan NN, <b>Huang CC</b>, 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.</li> <li>5. Huang CS, Lu TP, Liu CY, Huang CJ, Chiu JH, Chen YJ, Tseng LM, <b>Huang CC</b>. 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.</li> </ol>			
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