

Thesis Abstract

Functional analysis of pollutant nicotine-*CHRNA1* and identification of prognosis-related genes in melanoma using zebrafish and *in silico* approaches

(ゼブラフィッシュとインシリコ系を用いた、メラノーマにおけるニコチン - ニコチン性アセチルコリン受容体 A1 の機能解析と予後関連遺伝子の同定)

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Content of the thesis

Chapter 1.	General Introduction	5
	References	8
Chapter 2.	<i>CHRNA1</i> and its correlated-myogenesis/muscle contraction genes are prognosis-related markers of metastatic melanoma	12
	Introduction	13
	Material and Methods	15
	Results	22
	Discussion	28
	References	31
	Figures	37
	Supplementary Figures	44
	Tables	49
Chapter 3.	Analysis of Melanoma Gene Expression Signatures at the Single-Cell Level Uncovers 45-Gene Signature Related to Prognosis	50
	Introduction	51
	Material and Methods	54
	Results	59
	Discussion	66
	References	71
	Figures	82
	Supplementary Figures	94
	Supplementary Tables	99
Chapter 4.	General Discussion	101
	References	107
Acknowledgement	114