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Supplemental Information

Definition of miRNA Signatures of Nodal Metastasis in LCa: miR-449a Targets Notch Genes and Suppresses Cell Migration and Invasion

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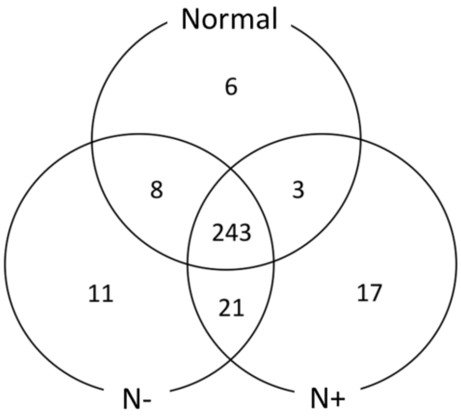
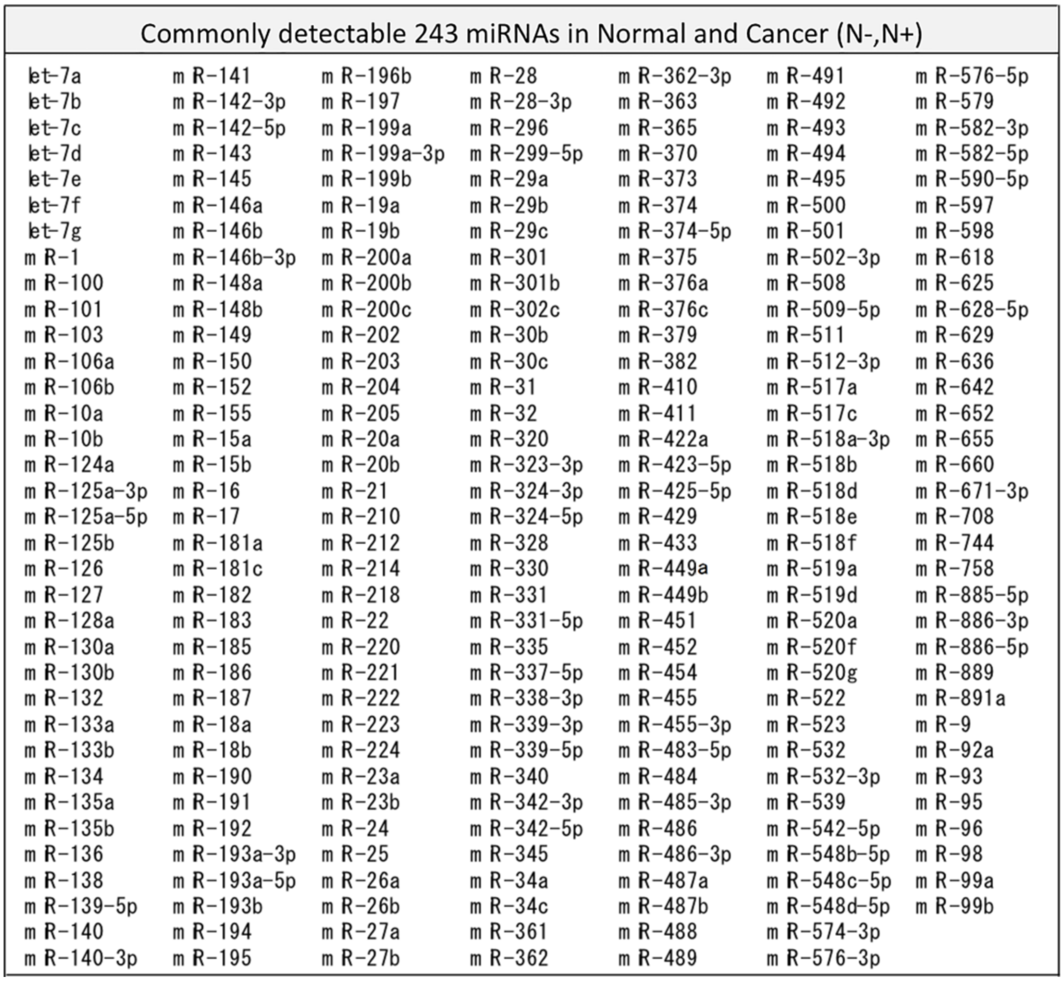
1A



1B


Commonly detectable 243 miRNAs in Normal and Cancer (N-,N+)

let-7a
let-7b
let-7c
let-7d
let-7e
let-7f
let-7g
miR-1
miR-100
miR-101
miR-103
miR-106a
miR-106b
miR-10a
miR-10b
miR-124a
miR-125a-3p
miR-125a-5p
miR-125b
miR-126
miR-127
miR-128a
miR-130a
miR-130b
miR-132
miR-133a
miR-133b
miR-134
miR-135a
miR-135b
miR-136
miR-138
miR-139-5p
miR-140
miR-140-3p
miR-141
miR-142-3p
miR-142-5p
miR-143
miR-145
miR-146a
miR-146b
miR-146b-3p
miR-148a
miR-148b
miR-149
miR-150
miR-152
miR-155
miR-15a
miR-15b
miR-16
miR-17
miR-181a
miR-181c
miR-182
miR-183
miR-185
miR-186
miR-187
miR-18a
miR-18b
miR-190
miR-191
miR-192
miR-193a-3p
miR-193a-5p
miR-193b
miR-194
miR-195
miR-196b
miR-197
miR-199a
miR-199a-3p
miR-199b
miR-19a
miR-19b
miR-200a
miR-200b
miR-200c
miR-202
miR-203
miR-204
miR-205
miR-20a
miR-20b
miR-21
miR-210
miR-212
miR-214
miR-218
miR-22
miR-220
miR-221
miR-222
miR-223
miR-224
miR-23a
miR-23b
miR-24
miR-25
miR-26a
miR-26b
miR-27a
miR-27b
miR-28
miR-28-3p
miR-296
miR-299-5p
miR-29a
miR-29b
miR-29c
miR-301
miR-301b
miR-302c
miR-30b
miR-30c
miR-31
miR-32
miR-320
miR-323-3p
miR-324-3p
miR-324-5p
miR-328
miR-330
miR-331
miR-331-5p
miR-335
miR-337-5p
miR-338-3p
miR-339-3p
miR-339-5p
miR-340
miR-342-3p
miR-342-5p
miR-345
miR-34a
miR-34c
miR-361
miR-362
miR-362-3p
miR-363
miR-365
miR-370
miR-373
miR-374
miR-374-5p
miR-375
miR-376a
miR-376c
miR-379
miR-382
miR-410
miR-411
miR-422a
miR-423-5p
miR-425-5p
miR-429
miR-433
miR-449a
miR-449b
miR-451
miR-452
miR-454
miR-455
miR-455-3p
miR-483-5p
miR-484
miR-485-3p
miR-486
miR-486-3p
miR-487a
miR-487b
miR-488
miR-489
miR-491
miR-492
miR-493
miR-494
miR-495
miR-500
miR-501
miR-502-3p
miR-508
miR-509-5p
miR-511
miR-512-3p
miR-517a
miR-517c
miR-518a-3p
miR-518b
miR-518d
miR-518e
miR-518f
miR-519a
miR-519d
miR-520a
miR-520f
miR-520g
miR-522
miR-523
miR-532
miR-532-3p
miR-539
miR-542-5p
miR-548b-5p
miR-548c-5p
miR-548d-5p
miR-574-3p
miR-576-3p
miR-576-5p
miR-579
miR-582-3p
miR-582-5p
miR-590-5p
miR-597
miR-598
miR-618
miR-625
miR-628-5p
miR-629
miR-636
miR-642
miR-652
miR-655
miR-660
miR-671-3p
miR-708
miR-744
miR-758
miR-885-5p
miR-886-3p
miR-886-5p
miR-889
miR-891a
miR-9
miR-92a
miR-93
miR-95
miR-96
miR-98
miR-99a
miR-99b

1C

Detectable 8 miRNAs in Normal and N-
miR-219-2-3p
miR-346
miR-369-5p
miR-383
miR-409-5p
miR-502
miR-627
miR-654-3p

Detectable 3 miRNAs in Normal and N+
miR-499
miR-545
miR-874

Detectable 21 miRNAs in N- and N+
miR-105
miR-129-3p
miR-147b
miR-184
miR-198
miR-296-3p
miR-424
miR-431
miR-450b-5p
miR-490
miR-501-3p
miR-503
miR-519c
miR-525-3p
miR-542-3p
miR-548d
miR-570
miR-589
miR-651
miR-873
miR-876-5p

Detectable 6 miRNAs in only Normal
miR-215
miR-450b-3p
miR-504
miR-211
miR-139-3p
miR-505

Detectable 11 miRNAs in only N-
miR-137
miR-216b
miR-372
miR-450a
miR-506
miR-510
miR-515-3p
miR-521
miR-654
miR-672
miR-674

Detectable 17 miRNAs in only N+
miR-107
miR-153
miR-208b
miR-369-3p
miR-381
miR-515-5p
miR-516b
miR-519e
miR-520a#
miR-551b
miR-872
miR-876-3p
miR-219-1-3p
miR-325
miR-517b
miR-520b
miR-888

Figure S1. Classification of detectable miRNAs among Normal, N- and N+ group.

(A) 309 miRNA candidates, detected at least 1 pool within 15 pool in PCR array assays, were selected and classified into 7 group of expression pattern by Venn diagram. (B) A commonly detected miRNA profile among Normal and Cancer (N-, N+) groups. (C) Detectable miRNAs in 2 different groups and solely miRNA expression profiles among Normal, N- and N+ group.

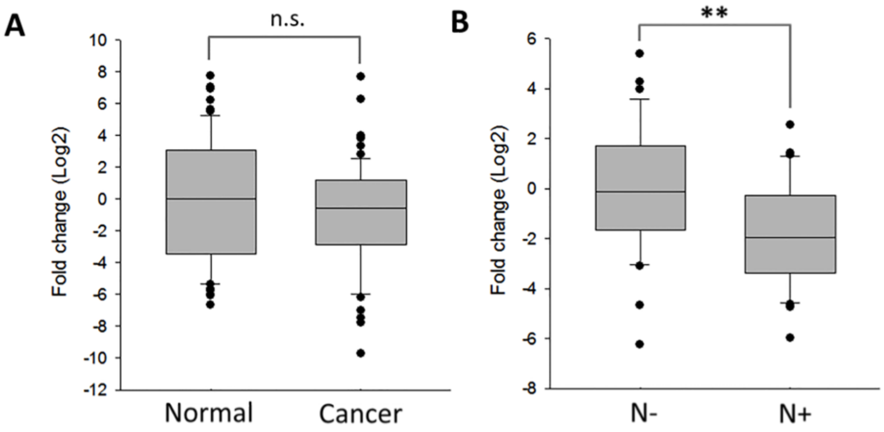


Figure S2. Validation of miR-449a expression in LCa tissues.

(A) The expression level of miR-449a was validated in LCa tissues with N+ (n = 32) and N- (n = 32) in a total of 64 LCa samples and their paired normal tissues (n = 64) using qRT-PCR. (B) miR-449a level in N+ (n = 32) compared to N- (n = 32). All results are shown as the box-and-whisker plot. For the normalization, U6 snRNA was used as endogenous control. Each sample was run in triplicate. Error bars show mean ± SD. The t test was used for the calculation of p-value. $*p < 0.05$, $**p < 0.01$.