

1 **Legends to figures and Figures**

2 **Supplementary figure 1.** Interaction sizes of target genes among five urinary miRNAs (miR-16-5p, miR-29a-3p, miR-124-3p, miR-199a-3p,
3 and miR-355-3p).

4 **Supplementary figure 2.** ROC analysis of urinary miRNA expression in distinguishing “progressors” among patients with IgAN. Each panel
5 represents ROC analysis: **a** miR-124-3p*miR-199a-3p*miR-355-3p*IIgANRPT, **b** miR-29a-3p* miR-124-3p*miR-199a-3p*miR-355-
6 3p*IIgANRPT, **c** miR-124-3p*miR-199a-3p*miR-355-3p, and **d** miR-29a-3p* miR-124-3p*miR-199a-3p*miR-355-3p).

7 **Supplementary figure 3.** Network of enrichment analysis of miR-124-3p.

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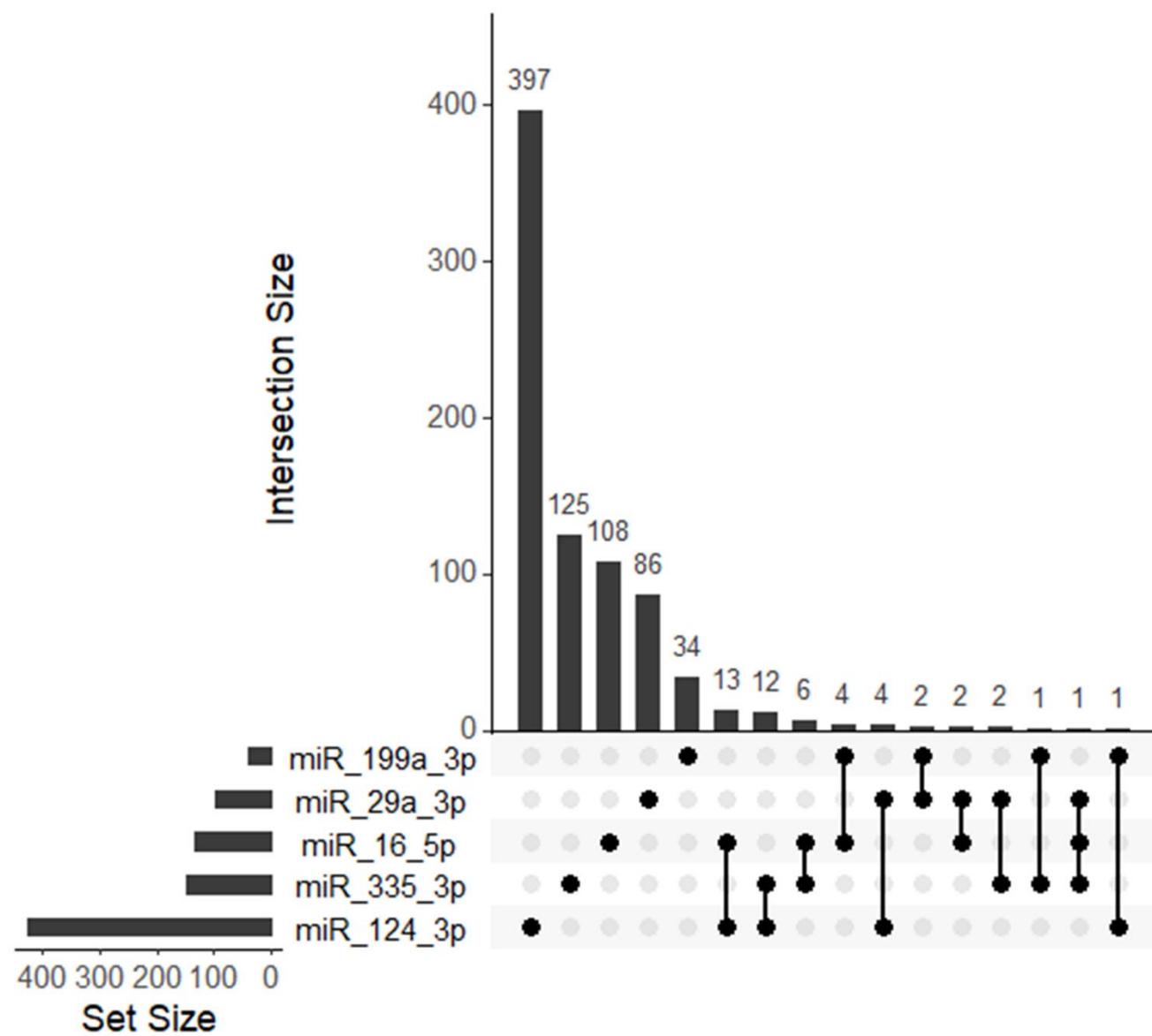
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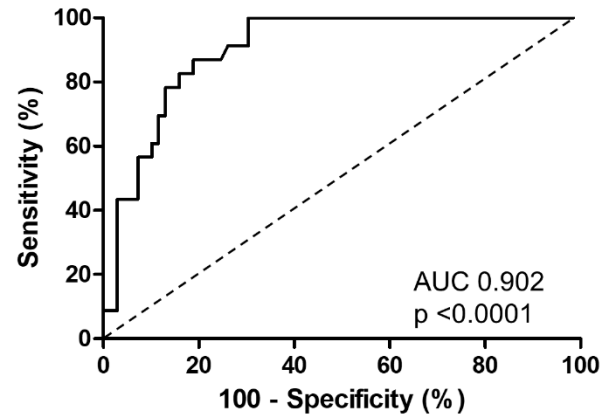
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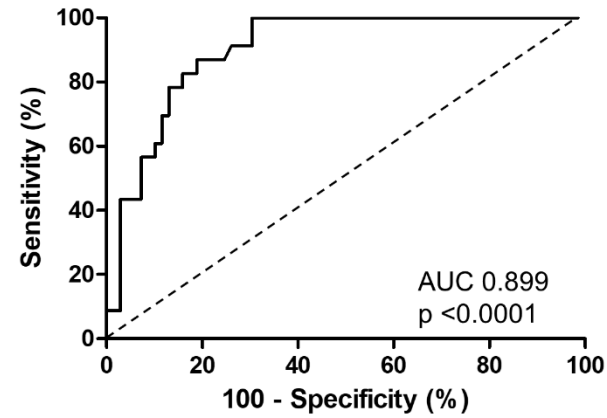
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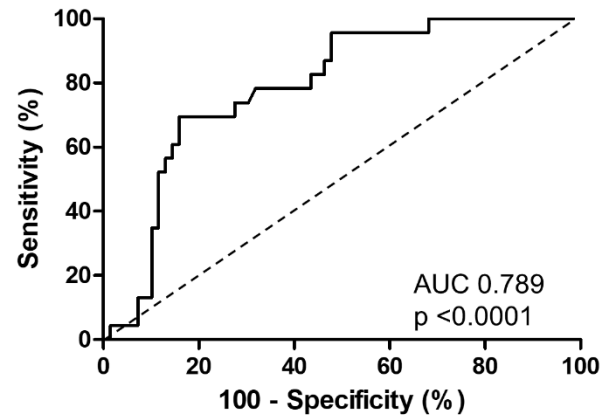
(A) miR-124-3p*miR-199a-3p*miR-335-3p*IlgANRPT



(B) miR-29a-3p*miR-124-3p*miR-199a-3p*miR-335-3p*IlgANRPT



(C) miR-124-3p*miR-199a-3p*miR-335-3p



(D) miR-29a-3p*miR-124-3p*miR-199a-3p*miR-335-3p

