

# The Pathophysiological Clarification Based on Inwardly Rectifying K Channel and the Development of Newly Antihypertensive Treatment in Salt Sensitive Hypertension

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## Summary

**Objectives:** Some aldosterone-producing adenoma (APA) has somatic mutation in *KCNJ5* coding for inwardly rectifying K channel (Kir3) which is mediated by G protein-coupled receptors (GPCRs). We aimed to detect novel genes associated with GPCRs in APA, and elucidate the mechanisms underlying aldosterone production.

**Methods:** Microarray analysis targeting GPCR-associated genes was conducted using APA without known mutations (APA-NM) samples ( $n = 8$ ) and APA samples with the *KCNJ5* mutation (APA-*KCNJ5*;  $n = 6$ ). Since gonadotropin-releasing hormone receptor (*GNRHR*) was one of the highest expression in APA-NM by microarray analysis, we investigated the effects of gonadotropin-releasing hormone (GnRH) stimulation on aldosterone production.

**Results:** Expression levels of mRNAs encoding *GNRHR* were highest in APA-NM samples according to our microarray analysis. The quantitative polymerase chain reaction (qPCR) assay results revealed higher *GNRHR* expression levels in APA-NM samples than in APA-*KCNJ5* samples ( $P < 0.05$ ). There was a significant and positive correlation between *GNRHR* expression and aldosterone increase via GnRH stimulation according to univariate and multivariate analyses. Consistent with the correlation, patients with APA-NM ( $n = 9$ ), which showed *GNRHR* mRNA levels, had significantly higher GnRH-stimulated aldosterone response than those with APA-*KCNJ5* ( $n = 13$ ) ( $P < 0.05$ ). We observed an aldosterone response in 55.6% (5/9) of patients with APA-NM, while none of the APA-*KCNJ5* patients exhibited an aldosterone response. A partial aldosterone response was seen in 22.2% (2/9) and 23.1% (3/13) of APA-NM and APA-*KCNJ5* patients, respectively. Multiple regression analysis revealed that the presence of the *KCNJ5* mutation was linked to *GNRHR* mRNA expression ( $\beta = 0.94$  and  $P < 0.01$ ). HAC15 cells with *KCNJ5* gene carrying T158A mutation exhibited a 0.64-fold increase in *GNRHR* expression than that in control cells ( $P < 0.05$ ).

**Conclusions:** We clarified increased expression of *GNRHR* in APA-NM, and the expression positively correlated with aldosterone production mediated by GnRH stimulation. Aberrant *GNRHR* expression in APA-NM could be one of the mechanisms by which aldosterone production is modulated.