LETTER TO THE EDITOR

GNAS Mutation Affecting Codon 201 is Rare in Most Human Tumors

Eun Mi Je • Chang Hyeok An • Yeun Jun Chung • Nam Jin Yoo • Sug Hyung Lee

Received: 15 September 2013 / Accepted: 19 February 2015 / Published online: 5 March 2015 © Arányi Lajos Foundation 2015

Keywords GNAS · Mutation · Cancers

To the Editors:

Guanine Nucleotide-binding protein, Alpha-Stimulating activity polypeptide 1 (GNAS) is a subunit of the stimulatory G protein that transmits signals to generate cAMP [1]. Mutations in GNAS gene result in several inherited diseases, including pseudohypoparathyroidism, pseudopseudohypoparathyroidism, progressive osseous heteroplasia, McCune-Albright syndrome, polyostotic fibrous dysplasia and several endocrine tumors [2]. Also, somatic mutations of GNAS have been reported in many tumors, including pituitary adenoma, colorectal carcinoma/adenoma, thyroid adenoma, juvenile ovarian granulosa cell tumor, low-grade appendix mucinous neoplasm, hepatocellular carcinomas, pancreatic mucinous neoplasm, fluke-associated cholangiocarcinoma and myelodysplasia [2-9]. Of note, most of the somatic GNAS mutations affected codon 201 [3-9]. Functionally, tumor-derived GNAS mutant constitutively activates cAMP signaling [3]. These data indicate that somatic mutations

E. M. Je · N. J. Yoo · S. H. Lee (⊠) Department of Pathology, College of Medicine, The Catholic University of Korea, 505 Banpo-dong, Socho-gu, Seoul 137-701, Korea e-mail: suhulee@catholic.ac.kr

C. H. An

Department of General Surgery, College of Medicine, The Catholic University of Korea, 505 Banpo-dong, Socho-gu, Seoul 137-701, Korea

Y. J. Chung

of *GNAS* are detected not only in endocrine tumors, but also in non-endocrine tumors [2–9].

To see whether the recurrent GNAS mutations occur in other tumor types as well, we analyzed the recurrent mutation site in 1, 126 tumor tissues (Table 1) by polymerase chain reaction (PCR) - single strand conformation polymorphism (SSCP) assay. All of the patients of the cancers were Asians (Koreans). We included myelodysplasia in this study that had been analyzed already, because in Korean population incidence of myelodysplasia subtypes were reported to be different from those of western countries. For example, incidence of MDS without RS is far less common than that of MDS with RS in Korean population [10]. In solid tumors, malignant cells and normal cells were selectively procured from hematoxylin and eosin-stained slides using a microdissection method. We analyzed exon 8 that encompassed the most common GNAS mutation site (codon 201). The primer sequences for amplification were as follows (forward and reverse, respectively): 5'-CTGTTTCGGTTGGCTTTGGTG -3' and 5'- AGGGACTGGGGTGAATGTCAAG -3'. SSCP analysis with this primer pair was proven to detect GNAS codon 201 mutation in our previous study [11]. Genomic DNA each from tumor cells and corresponding normal cells were amplified with the primer pairs by PCR. Radioisotope was incorporated into the PCR products for detection by autoradiogram.

However, SSCPs from the 1,126 cancers did not reveal any aberrantly migrating band compared to bands from normal tissues, indicating there was no evidence of mutation in the recurrent mutation sites in *GNAS* gene in the tumors. In a positive control (intraductal papillary mucinous neoplasm of the pancreas), the SSCP detected aberrantly migrating bands. We repeated the experiments twice, including tissue microdissection, PCR and SSCP analysis to ensure the results, and found that the data was consistent.

Prior to this study, we expected to find the recurrent mutations of GNAS in our samples analyzed, because GNAS recurrent mutations have been detected in many tumor types [3–9] and our

Department of Microbiology, College of Medicine, The Catholic University of Korea, 505 Banpo-dong, Socho-gu, Seoul 137-701, Korea

Table 1 Tumor tissues analyzed in this study

Type of tumors	Number of tumor
Head/neck squamous carcinoma	13
Esophageal squamous carcinoma	59
Ovarian carcinoma	54
Prostate carcinoma	267
Non-malanomaous skin cancers	26
Hepatoblastoma	14
Sarcomas	64
Uterine leiomyoma	68
Meningioma	13
Gastrointestinal stromal tumor	19
Ovarian stromal tumor	43
Acute myelogenous leukemia	180
Acute lymphoblastic leukemia	180
Myelodysplasia	68
Non-Hodgkin lymphoma	58
Total	1,126

cancer samples included diverse cancer types (Table 1). However, we detected no GNAS mutations in those cancers (Table 1). Moreover, there was no GNAS mutation in myelodysplasia where the GNAS mutation had been detected [8]. Comparing the incidence of GNAS mutation in myelodysplasia between our data (0/68) and the earlier data (3/439), we found no significant difference between the two (Fisher's exact test, p=0.649), indicating that the recurrent GNAS mutation may be a rare event in myelodysplasia. In non-endocrine tumors, high incidences of the GNAS mutations have been reported in papillary mucinous neoplasm of digestive system [4, 7]. In other tumors, the incidences were negative or very low [5, 10]. Together, the data suggest that recurrent GNAS mutations are present in special types of tumors (endocrine tumors and papillary mucinous neoplasm), but rare in other tumors.

In summary, our study did not find any *GNAS* recurrent mutation in carcinomas (head/neck, esophagus, ovary, skin, liver and prostate), stromal tumors (sarcomas, leiomyomas, meningiomas, ovarian stromal tumors and gastrointestinal stromal tumors) and hematologic tumors (leukemias, myelodysplasia and lymphomas), indicating that *GNAS* codon 201 may be rarely mutated in most tumor types.

Acknowledgments This study was supported by a grant from Korea Research Foundation (2012R1A5A20047939).

References

- Landis CA, Masters SB, Spada A, Pace AM, Bourne HR, Vallar L (1989) GTPase inhibiting mutations activate the alpha chain of Gs and stimulate adenylyl cyclase in human pituitary tumours. Nature 340:692–696
- 2. Aldred MA, Trembath RC (2000) Activating and inactivating mutations in the human GNAS1 gene. Hum Mutat 16:183–189
- 3. Sjöblom T, Jones S, Wood LD, Parsons DW, Lin J, Barber TD, Mandelker D, Leary RJ, Ptak J, Silliman N, Szabo S, Buckhaults P, Farrell C, Meeh P, Markowitz SD, Willis J, Dawson D, Willson JK, Gazdar AF, Hartigan J, Wu L, Liu C, Parmigiani G, Park BH, Bachman KE, Papadopoulos N, Vogelstein B, Kinzler KW, Velculescu VE (2006) The consensus coding sequences of human breast and colorectal cancers. Science 314:268–274
- Nishikawa G, Sekine S, Ogawa R, Matsubara A, Mori T, Taniguchi H, Kushima R, Hiraoka N, Tsuta K, Tsuda H, Kanai Y (2013) Frequent GNAS mutations in low-grade appendiceal mucinous neoplasms. Br J Cancer 108:951–958
- Nault JC, Fabre M, Couchy G, Pilati C, Jeannot E, Tran Van Nhieu J, Saint-Paul MC, De Muret A, Redon MJ, Buffet C, Salenave S, Balabaud C, Prevot S, Labrune P, Bioulac-Sage P, Scoazec JY, Chanson P, Zucman-Rossi J (2012) GNAS-activating mutations define a rare subgroup of inflammatory liver tumors characterized by STAT3 activation. J Hepatol 56:184–191
- Kalfa N, Lumbroso S, Boulle N, Guiter J, Soustelle L, Costa P, Chapuis H, Baldet P, Sultan C (2006) Activating mutations of Gsalpha in kidney cancer. J Urol 176:891–895
- Furukawa T, Kuboki Y, Tanji E, Yoshida S, Hatori T, Yamamoto M, Shibata N, Shimizu K, Kamatani N, Shiratori K (2011) Wholeexome sequencing uncovers frequent GNAS mutations in intraductal papillary mucinous neoplasms of the pancreas. Sci Rep 1:161
- Bejar R, Stevenson K, Abdel-Wahab O, Galili N, Nilsson B, Garcia-Manero G, Kantarjian H, Raza A, Levine RL, Neuberg D, Ebert BL (2011) Clinical effect of point mutations in myelodysplastic syndromes. N Engl J Med 364:2496–2506
- Ong CK, Subimerb C, Pairojkul C, Wongkham S, Cutcutache I, Yu W, McPherson JR, Allen GE, Ng CC, Wong BH, Myint SS, Rajasegaran V, Heng HL, Gan A, Zang ZJ, Wu Y, Wu J, Lee MH, Huang D, Ong P, Chan-on W, Cao Y, Qian CN, Lim KH, Ooi A, Dykema K, Furge K, Kukongviriyapan V, Sripa B, Wongkham C, Yongvanit P, Futreal PA, Bhudhisawasdi V, Rozen S, Tan P, Teh BT (2012) Exome sequencing of liver fluke-associated cholangiocarcinoma. Nat Genet 44:690–693
- Jung SW, Lee SY, Jekarl DW, Kim M, Lim J, Kim Y, Han K, Kim YJ, Cho SG, Song J (2011) Cytogenetic characteristics and prognosis analysis in 231 myelodysplastic syndrome patients from a single institution. Leuk Res 35:735–740
- Lee SH, Jeong EG, Soung YH, Lee JW, Yoo NJ, Lee SH (2008) Absence of GNAS and EGFL6 mutations in common human cancers. Pathology 40:95–97