

Supplementary Table. Characteristics of allele frequency of the identified mutations

Patients who fulfill NIFTP 2016 and NIFTP 2018 diagnostic criteria				
Case number	Gene	Mutation position	AA change	Allele frequency (%)
1	<i>HRAS</i>	c.A182G	p.Q61R	13
2	<i>TP53</i>	c.G569A	p.C190Y	5
	<i>SMAD4</i>	G302A	p.W101X	8
3	<i>NRAS</i>	c.A182G	p.Q61R	42
4	<i>HRAS</i>	c.C100T	p.P34S	48
	<i>HNF1A</i>	c.G599A	p.R200Q	37
	<i>TP53</i>	c.C457T	p.Q153X	31
5	<i>MET</i>	c.A3758G	p.Y1253C	5
6	<i>KIT</i>	c.G1681A	p.G561R	4
7	<i>KRAS</i>	c.A182G	p.Q61R	4
	<i>APC</i>	c.C4348T	p.R1450X	11
8	<i>TP53</i>	c.G701T	p.R234L	4
9	<i>KRAS</i>	c.A182G	p.Q61R	8
10	<i>ATM</i>	c.A1052T	p.D351V	34
11	<i>SMARCB1</i>	c.G152A	p.W51X	6
	<i>KIT</i>	c.G1576A	p.V526I	34
12	<i>PTEN</i>	c.C1030T	p.Q344X	7
13	<i>KRAS</i>	c.G40A	p.V14I	4
Patients who fulfill only NIFTP 2016 diagnostic criteria				
Case number	Genetic mutations	Mutation position	AA change	Allele frequency (%)
1	<i>BRAF</i>	c.T1799A	p.V600E	4
2	<i>BRAF</i>	c.T1799A	p.V600E	6
3	<i>BRAF</i>	c.T1799A	p.V600E	7
4	<i>BRAF</i>	c.T1799A	p.V600E	35
5	<i>BRAF</i>	c.T1799A	p.V600E	39
6	<i>BRAF</i>	c.1796_1797insTAC	p.T599delinsTT	31
7	<i>TP53</i>	c.A619T	p.M207L	10
8	<i>APC</i>	c.C4073T	p.A1358V	50
9	<i>KRAS</i>	c.A182G	p.Q61R	4
10	<i>KRAS</i>	c.G34C	p.G12R	12

NIFTP, noninvasive follicular thyroid neoplasm with papillary-like nuclear features; AA, amino acids