

SUPPLEMENTAL

Table S1. Details of genetic variants on *EIF2AK4* gene

| No. | ID | Gender | Gene | cDNA change | Amino acid change | Variant type | Genotype | PolyPhen2 /SIFT | ACMG 2015* | MAF in EAS | Ref |
|-----|---------|--------|----------------|-------------|-------------------|--------------|----------|-----------------|------------|------------|-----|
| | A100540 | M | <i>EIF2AK4</i> | c.4218G>T | p.Gln1406His | Missense | Het | B/T | LB | 0.01177 | |
| | A100551 | M | <i>EIF2AK4</i> | c.4218G>T | p.Gln1406His | Missense | Het | B/T | LB | 0.01177 | |
| | A100559 | F | <i>EIF2AK4</i> | c.4218G>T | p.Gln1406His | Missense | Het | B/T | LB | 0.01177 | |
| | A100641 | F | <i>EIF2AK4</i> | c.4218G>T | p.Gln1406His | Missense | Het | B/T | LB | 0.01177 | |
| | A110137 | F | <i>EIF2AK4</i> | c.4218G>T | p.Gln1406His | Missense | Het | B/T | LB | 0.01177 | |

1. Reference sequences: *EIF2AK4* (NM_001013703.4)

2. PolyPhen2: PD, Probably damaging; D, Damaging; B, Benign. SIFT: D, Deleterious, T, Tolerated.

3. PolyPhen2 and SIFT *in silico* prediction program were unable to analyze frameshift, nonsense, in frame deletion, gross deletion types and some variants with unknown reasons.

4. ACMG 2015: The American College of Medical Genetics and Genomics guidelines : VUS, variant of uncertain significance; LB, Likely Benign.

5. MAF in EAS: Minor allele frequency of East Asian in gnomAD exome databases.

Table S2. Female pulmonary arterial hypertension patients with or without bone morphogenetic protein receptor type 2 (*BMPR2*) genetic variants, clinical and hemodynamic presentations at initial diagnosis (N=51).

| Characteristics | | <i>BMPR2</i> variant carrier (N=5) | Non- <i>BMPR2</i> , non- <i>GDF2</i> variant (N=46) | <i>p</i> -value <i>BMPR2</i> versus non- <i>BMPR2</i> variant |
|---|-----|---|--|--|
| Female (N, %)) | | 5 (100) | 46 (100) | |
| Age of onset, years | | 43 ± 11 | 54 ± 20.2 | 0.277 |
| Six-minute walking distance (m) | | 351 ± 165 | 322 ± 123 | 0.613 |
| Mean pulmonary arterial pressure (mmHg) | | 66 ± 15 | 37 ± 14 | <0.001 |
| Pulmonary arterial wedge pressure (mmHg) | | 11 ± 3 | 14 ± 5 | 0.198 |
| Pulmonary vascular resistance (woods) | | 22 ± 9 | 7 ± 5 | <0.001 |
| Peak tricuspid regurgitation peak gradient (mmHg) | | 64 ± 35 | 49 ± 31 | 0.330 |
| Peak oxygen consumption (mL/min/kg) | | 12 ± 4 | 12 ± 4 | 0.948 |
| ventilatory equivalents for carbon dioxide (VE/VCO2) | | 50 ± 18 | 38 ± 13 | 0.088 |
| Right atrial pressure (mmHg) | | 14 ± 9 | 14 ± 12 | 0.994 |
| Cardiac index (L/min/m²) | | 1.6 ± 0.5 | 2.7 ± 1.1 | 0.021 |
| Pulmonary artery saturation (%) | | 54 ± 13 | 65 ± 12 | 0.060 |
| N-terminal prohormone of brain natriuretic peptide (ng/L) | | 4117 ± 533 6 | 1875 ± 2897 | 0.139 |
| World Health Organization functional class (N, %) | I | 1 (20) | 3 (6.5) | 0.490 |
| | II | 1 (20) | 11 (23.9) | |
| | III | 3 (60) | 30 (65.2) | |
| | IV | 0 (0) | 2 (4.3) | |
| Progression of symptoms | No | 1 (20) | 19 (41.3) | 0.502 |

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| Slow | 4 (80) | 21 (45.7) |
| Rapid | 0 (0) | 6 (13.0) |

† Data of continuous variables were expressed as mean \pm SD.

† Changes of categorical variables were analyzed by Chi-square tests and were expressed by (N, %).

Table S3. Male Pulmonary arterial hypertension patients with or without Growth Differentiation Factor 2 (*GDF2*) genetic variants, clinical and hemodynamic presentations at initial diagnosis (N=18).

| Characteristics | <i>GDF2</i> variant carrier (N=3) | Non- <i>BMPR2</i> , non- <i>GDF2</i> variant (N=15) | <i>p</i> -value |
|---|-----------------------------------|---|---|
| | | | <i>GDF2</i> versus non- <i>GDF2</i> variant |
| Male | 3 (100) | 15 (100) | |
| Age of onset, years | 25 ± 13 | 44 ± 19 | 0.109 |
| Six-minute walking distance (m) | 490 ± 101 | 319 ± 151 | 0.082 |
| Mean pulmonary arterial pressure (mmHg) | 66 ± 13 | 39 ± 12 | 0.002 |
| Pulmonary arterial wedge pressure (mmHg) | 11 ± 3 | 14 ± 2 | 0.138 |
| Pulmonary vascular resistance (woods) | 15 ± 12 | 6 ± 5 | 0.048 |
| Peak tricuspid regurgitation peak gradient (mmHg) | 58 ± 23 | 54 ± 30 | 0.827 |
| Peak oxygen consumption (mL/min/kg) | 15 ± 6 | 14 ± 5 | 0.722 |
| ventilatory equivalents for carbon dioxide (VE/VCO ₂) | 45 ± 4 | 36 ± 17 | 0.500 |
| Right atrial pressure (mmHg) | 11 ± 5 | 12 ± 4 | 0.683 |
| Cardiac index (L/min/m ²) | 2.6 ± 1.1 | 2.9 ± 1.0 | 0.734 |
| Pulmonary artery saturation (%) | 68 ± 10 | 71 ± 8 | 0.612 |
| N-terminal prohormone of brain natriuretic peptide (ng/L) | 228 ± 46 | 1396 ± 2324 | 0.410 |
| World Health Organization functional class (N, %) I | 0 (0) | 5 (33.3) | 0.840 |
| II | 2 (66.7) | 8 (53.3) | |
| III | 1 (33.3) | 2 (13.3) | |
| IV | 0 (0) | 0 (3.3) | |
| Progression of symptoms No | 1 (33.3) | 5 (38.5) | 0.670 |

| | | | |
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| | Slow | 1 (33.3) | 8 (61.5) |
| | Rapid | 1 (33.3) | 2 (15.4) |

† Data of continuous variables were expressed as mean \pm SD.

† Changes of categorical variables were analyzed by Chi-square tests and were expressed by (N, %).

Table S4. Pulmonary arterial hypertension patients with or without *ATP13A3* genetic variants, clinical and hemodynamic presentations at initial diagnosis (N=63).

| Characteristics | | <i>ATP13A3</i> variant carriers (N=4) | Non- <i>ATP13A3</i> , variant carriers (N=59)* | <i>p</i> -value |
|---|------|---|--|-----------------|
| Female (N, %)) | | 3 (75) | 45 (76.3) | |
| Age of onset, years | | 48.0 ± 8.6 | 51.4 ± 20.4 | 0.746 |
| Six-minute walking distance (m) | | 320 ± 167 | 319 ± 122 | 0.982 |
| Mean pulmonary arterial pressure (mmHg) | | 52 ± 7 | 38 ± 13 | 0.043 |
| Pulmonary arterial wedge pressure (mmHg) | | 12 ± 2 | 14 ± 5 | 0.296 |
| Pulmonary vascular resistance (woods) | | 15 ± 14 | 7 ± 5 | 0.353 |
| Peak tricuspid regurgitation peak gradient (mmHg) | | 45 ± 29 | 51 ± 30 | 0.687 |
| Peak oxygen consumption (mL/min/kg) | | 11 ± 6 | 12 ± 4 | 0.528 |
| ventilatory equivalents for carbon dioxide (VE/VCO2) | | 44 ± 19 | 38 ± 14 | 0.406 |
| Right atrial pressure (mmHg) | | 13 ± 8 | 14 ± 10 | 0.881 |
| Cardiac index (L/min/m ²) | | 1.8 ± 0.8 | 2.8 ± 1.1 | 0.078 |
| Pulmonary artery saturation (%) | | 57 ± 12 | 67 ± 12 | 0.116 |
| N-terminal prohormone of brain natriuretic peptide (ng/L) | | 4254 ± 5876 | 1782 ± 2806 | 0.120 |
| World Health Organization functional class (N, %) | I | 0 (0) | 3 (5.1) | 0.656 |
| | II | 0 (0) | 16 (27.1) | |
| | III | 3 (100) | 38 (64.4) | |
| | IV | 0 (0) | 2 (3.4) | |
| | | | | |
| Progression of symptoms | No | 1 (25) | 23 (40.0) | 0.704 |
| | Slow | 2 (50) | 29 (49.2) | |

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|-------|----------|------------|
| Rapid | 1 (25) | 7 (11.9) |
|-------|----------|------------|

† Data of continuous variables were expressed as mean \pm SD.

† Changes of categorical variables were analyzed by Chi-square tests and were expressed by (N, %).

* Non-ATP13A3 carrier group also excluded patients carrying variants on *BMPR2* and *GDF2*.

Table S5 . Pulmonary arterial hypertension patients with or without *EIF2AK4* genetic variants, clinical and hemodynamic presentations at initial diagnosis (N=62).

| Characteristics | <i>EIF2AK4</i> variant carriers (N=9) | Non- <i>EIF2AK4</i> variant carriers (N=53)* | <i>p</i> -value |
|---|---|--|-----------------|
| Female (N, %)) | 3 (75) | 45 (76.3) | |
| Age of onset, years | 52.4 ± 23.6 | 50.4 ± 20.1 | 0.779 |
| Six-minute walking distance (m) | 334 ± 153 | 321 ± 117 | 0.767 |
| Mean pulmonary arterial pressure (mmHg) | 42 ± 12 | 38 ± 14 | 0.345 |
| Pulmonary arterial wedge pressure (mmHg) | 14 ± 6 | 14 ± 4 | 0.798 |
| Pulmonary vascular resistance (woods) | 6 ± 4 | 7 ± 5 | 0.470 |
| Peak tricuspid regurgitation peak gradient (mmHg) | 50 ± 43 | 50 ± 28 | 0.983 |
| Peak oxygen consumption (mL/min/kg) | 14 ± 4 | 12 ± 4 | 0.226 |
| ventilatory equivalents for carbon dioxide (VE/VCO2) | 36 ± 10 | 38 ± 15 | 0.744 |
| Right atrial pressure (mmHg) | 11 ± 5 | 14 ± 11 | 0.450 |
| Cardiac index (L/min/m ²) | 3.1 ± 1.3 | 2.7 ± 1.0 | 0.421 |
| Pulmonary artery saturation (%) | 66 ± 12 | 67 ± 11 | 0.951 |
| N-terminal prohormone of brain natriuretic peptide (ng/L) | 3526 ± 5117 | 1428 ± 2027 | 0.258 |
| World Health Organization functional class (N, %) | | | |
| I | 0 (0) | 3 (5.9) | 0.417 |
| II | 1 (11) | 16 (31.4) | |
| III | 8 (89) | 32 (62.7) | |
| IV | 0 (0) | 2 (3.9) | |
| Progression of symptoms | No | 3 (33.3) | 0.897 |

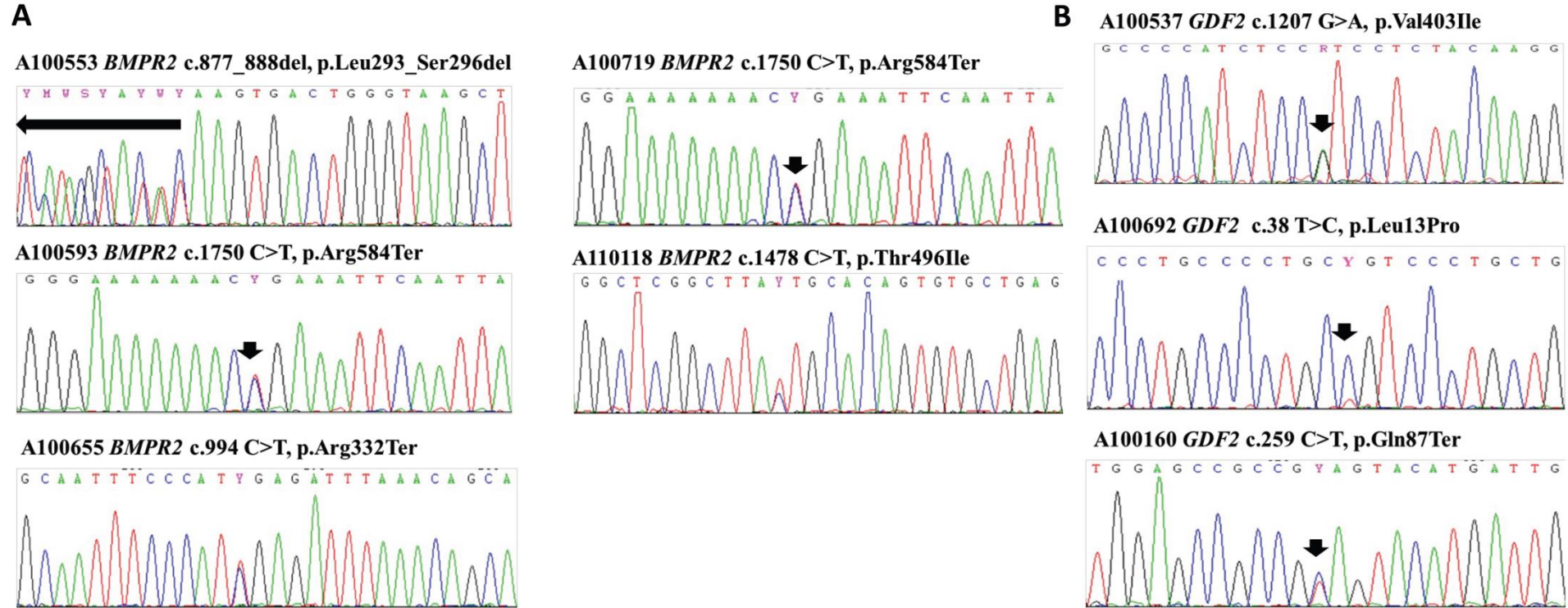
| | | |
|-------|------------|-------------|
| Slow | 5 (55.6) | 25 (47.2) |
| Rapid | 1 (11.1) | 7 (13.2) |

† Data of continuous variables were expressed as mean \pm SD.

† Changes of categorical variables were analyzed by Chi-square tests and were expressed by (N, %).

* Non-*EIF2AK4* carriers group also excluded patients carrying variants on *BMPR2* and *GDF2*.

Supplemental Figure 1.



Supplement Figure S1. Sanger sequencing validation of variants on (A) *BMPR2* and (B) *GDF2*. Five *BMPR2* variants and three *GDF2* variants from IPAH patients were confirmed by PCR-Sanger sequencing directly. Black arrows indicated exact variant. Y means C to T transition. R means A to G transition.