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Genetic Variation of SCNN1A Influences Lung Diffusing Capacity in Cystic Fibrosis

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Abstract

Introduction—Epithelial Na⁺ Channels (ENaC) play a crucial role in ion and fluid regulation in the lung. In cystic fibrosis (CF) Na⁺ hyperabsorption results from ENaC over activity, leading to airway dehydration. Previous work has demonstrated functional genetic variation of SCNN1A (the gene encoding the ENaC α -subunit), manifesting as an alanine (A) to threonine (T) substitution at amino acid 663, with the α T663 variant resulting in a more active channel.

Methods—We assessed the influence of genetic variation of SCNN1A on the diffusing capacity of the lungs for carbon monoxide (DLCO) and nitric oxide (DLNO), together with alveolar capillary membrane conductance (D_M), pulmonary capillary blood volume (V_C), and alveolar volume (V_A) at rest and during peak exercise in 18 patients with CF [10 homozygous for α A663 (AA group) and 8 with at least one T663 allele (AT/TT group)]. Due to the more active channel we hypothesized that the AT/TT group would show a greater increase in DLCO, DLNO, and D_M with exercise due to exercise-mediated ENaC inhibition and subsequent attenuation of Na⁺ hyperabsorption.

Results—The AT/TT group had significantly lower pulmonary function, weight and BMI than the AA group. Both groups had similar peak workloads, relative peak oxygen consumptions, and cardiopulmonary responses to exercise. The AT/TT group demonstrated a greater increase in DLNO, DLNO/ V_A , and D_M in response to exercise (% increases: DLNO= 18±11 vs.41±38; DLNO/ V_A = 14±21 vs.40±37; D_M = 15±11 vs.41±38, AA vs. AT/TT, respectively). There were no differences between groups in absolute diffusing capacity measures at peak exercise.

Conclusion—These results suggest that genetic variation of the alpha-subunit of ENaC differentially affects the diffusing capacity response to exercise in patients with CF.

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exercise; cystic fibrosis; diffusing capacity; DLNO; DLCO; ENaC polymorphism

Introduction

Cystic fibrosis (CF) is an autosomal recessive disorder caused by mutations in the gene encoding the cystic fibrosis transmembrane conductance regulator (CFTR) which affects 1 in 3419 white individuals and 1 in 12,163 non-white individuals (21). Cystic fibrosis transmembrane conductance regulator helps to maintain airway surface fluid depth through secretion of Cl^- ; the impairment in CFTR results in a depletion of the airway surface fluid. There are several mutations which result in CF, with the ΔF508 mutation being the most common (13). The ΔF508 mutation is a deletion of phenylalanine at amino acid 508 which causes defective trafficking of CFTR and results in the inability of CFTR to be inserted into the apical membrane of the cell (25).

Extravascular lung water has been estimated to be 3.6mL/Kg in healthy individuals, and maintenance of extravascular lung water is important for proper gas diffusion to occur (37). The diffusing capacity of the lung is reduced in disease states like CF where the lungs are dry, whereas if extravascular lung water increases, like during rapid saline loading or in heart failure, the diffusing capacity of the lung is also reduced (28, 30, 39). Airway surface fluid depth is of vital importance in pulmonary defense and function by aiding in mucus clearance in the airways. In the alveoli, the maintenance of optimal airway surface fluid depth is also essential in gas diffusion across the respiratory membrane (12). Increases in surface fluid depth are mediated primarily by Cl^- secretion by CFTR, but also occur through Cl^- secretion by Ca^{2+} -dependent Cl^- channels (CaCC) (5, 8). Surface fluid depth is decreased primarily through Na^+ reabsorption by the epithelial Na^+ channel (ENaC) (9, 11). Sodium is brought into the epithelial cell by ENaC where it is subsequently moved across the basolateral membrane by Na^+/K^+ ATPase into the interstitial space. Fluid follows through tight junctions and aquaporin channels where it is then cleared by the lymphatic system. Not only is Cl^- secretion impaired in CF, but Na^+ absorption via ENaC is dramatically increased due to the loss of CFTR mediated ENaC inhibition, potentiating the decreased surface liquid volume seen in CF, as well as impairing mucociliary clearance, leading to airway obstruction, inflammation, recurrent infections, and decreasing the diffusing capacity of gases across the respiratory membrane (39).

Previous work has shown that exercise increases the diffusing capacity of gases in the lungs of both healthy individuals and those with CF (39). Individuals with CF demonstrated lower diffusing capacity of the lungs at rest and during exercise as compared to healthy individuals (39). Exercise can activate ENaC through the adrenergic pathway but can also inhibit ENaC through the purinergic pathway, which is activated by turbulent air flow in the airways during exercise (1). Several functional polymorphisms in the gene encoding ENaC (SCNN1A) have been reported, including a common variant resulting in an alanine (A) to threonine (T) substitution at amino acid 663 in the C-terminus of the alpha subunit. The more common variant (αA663) demonstrated lower channel activity as compared to the αT663 variant due to decreased ENaC expression (27, 35). In humans, the αT663 variant has been associated with hypertension, suggesting its increased channel activity resulted in increased Na^+ reabsorption in the kidneys (2). Genetic variation of SCNN1A has also previously been shown to influence the diffusing capacity of the lungs in response to β_2 -adrenergic stimulation (3). Healthy individuals with the αA663 ENaC variant demonstrated a greater increase in lung diffusing capacity in response to exercise as compared to those with at least one copy of the αT663 variant (3). Additionally, in response to albuterol, an

exogenous β_2 -receptor agonist, individuals with the α A663 ENaC variant had less Na^+ in their exhaled breath condensate 90 minutes after β_2 -agonist administration as compared to those with the α T663 ENaC variant, suggesting a greater removal of Na^+ from the airways following activation (14).

In this study, we sought to determine the influence of genetic variation at amino acid 663 of the alpha-subunit of ENaC on diffusing capacity of the lungs at rest and during exercise in patients with CF. Exercise was utilized as a means to endogenously alter ion and fluid regulation in the lung. We predicted that patients with the α T663 ENaC variant, which has shown greater basal activity in previous work (27, 35), would show a greater increase in diffusing capacity with exercise, due to a larger attenuation of Na^+ reabsorption, and subsequent hydration of the airways.

Methods

Subjects

Eighteen subjects with mild to moderate CF, with a positive sweat test ($>60\text{mmol/L}$) and at least one ΔF508 allele, were recruited for study through the Arizona Respiratory Center and its affiliated CF clinic at the University of Arizona Medical Center. To ensure subjects were clinically stable to participate in the study exclusion criteria included: $\text{FEV}_1 < 40\%$ predicted, experimental CF drugs, pulmonary exacerbation within the last six months resulting in $>50\text{cc}$ of blood in the sputum, antibiotics for pulmonary exacerbation, pregnancy, smoking, inability to exercise, or cardiovascular abnormalities. The protocol was reviewed and approved by the University of Arizona Institutional Review Board, and all subjects provided written informed consent before participation in the study. All aspects of the study were performed according to the declaration of Helinski.

Protocol

Upon arrival to the laboratory in a two hour fasted state, subjects were consented and a buccal swab was performed for assessment of the amino acid at position 663 of αENaC and verification of ΔF508 mutation on CFTR. A venous blood sample was drawn for the assessment of hemoglobin concentration. Subjects were outfitted with a 12-lead electrocardiogram (Marquette electronics, Milwaukee, WI) to monitor heart rhythms, and pulmonary function tests were completed according to American Thoracic Society standards (23). Predicted values for pulmonary function tests were calculated according to equations from NHANES III (17). Baseline simultaneous measurements of the diffusing capacity of the lungs for carbon monoxide (DLCO) and the diffusing capacity of the lungs for nitric oxide (DLNO) were taken in triplicate. Baseline peripheral oxygen saturation (SaO_2) was assessed via pulse oximetry with a finger sensor (Nellcor N-600 Pulse Oximeter, Boulder, CO). Subjects then completed a subject-specific maximal exercise capacity test on a cycle ergometer (Corival Lode B.V., The Netherlands) based on their body size, reported type, speed and intensity of exercise training, and predicted \dot{V}_{O_2} (18). DLCO and DLNO were taken during each stage of exercise and recovery. Subjects exercised at an initial workload that ranged from 15-40 watts (mean initial workload was 24 ± 6 watts) and the workload was increased by this same amount every three minutes until exhaustion (ie. initial workload of 25 watts with a 25 watt increase in workload every three minutes). The exercise test ended with a three-minute recovery period at the initial workload. Exhaustion was determined by an inability to maintain a pedal rate of 60-80 revolutions per minute, a rating of perceived exertion (RPE) of at least 18 out of 20, or a respiratory exchange ratio (RER) greater than or equal to 1.15.

Oxygen consumption (\dot{V}_{O_2}), carbon dioxide production (\dot{V}_{CO_2}), respiratory rate (RR), tidal volume (T_V), minute ventilation (V_E), heart rate (HR), and oxygen saturation (SaO_2) were monitored continuously and averaged every three seconds throughout the test via a metabolic cart (Medical Graphics CPX/D; St. Paul, MN) interfaced with a Perkin Elmer MGA-1100 mass spectrometer (Wesley, MA) as described previously (30, 31). Accurate SaO_2 values were ensured by instructing the subjects to maintain a relaxed grip and verifying that there were no discrepancies in HR between the pulse oximeter HR and that on the electrocardiogram.

Measurement of cardiac output, alveolar-capillary membrane conductance, and pulmonary capillary blood volume

Cardiac output, pulmonary capillary blood volume (V_C) and alveolar-capillary membrane conductance (D_M) were determined using a rebreath technique by measuring the disappearance of acetylene, carbon monoxide (CO), and nitric oxide (NO) with respect to helium, as previously described (16, 30-33). Triplicate maneuvers of cardiac output (Q), the diffusing capacity of the lungs for carbon monoxide (DLCO), and nitric oxide (DLNO) were performed at baseline and during recovery. Single maneuvers of Q, DLCO, and DLNO were performed during each stage of exercise.

Measures of Q, DLCO, and DLNO were taken in an upright and seated position on the cycle ergometer using a rebreathing technique, with gases sampled by a Perkin Elmer MGA-1100 mass spectrometer (Wesley, MA) and Seivers Instruments NO analyzer (Boulder, CO) integrated with custom analysis software, as described previously (29, 31, 38, 39). Subjects breathed into a five-liter rebreathing bag containing 9% helium, 0.7% acetylene, 0.3% carbon monoxide ($C^{18}O$), 40 PPM NO, and 35% O_2 with a respiratory rate controlled at 32 breaths per minute via metronome. The 40 PPM NO was diluted in the anesthesia bag immediately prior to each maneuver from an 800 PPM NO tank. $C^{18}O$ was used instead of the more common $C^{16}O$ to enable the mass spectrometer to distinguish between $C^{16}O$ and N_2 , which have similar molecular weights. The total volume of gas used in the rebreathing maneuver was standardized at 1575 mL at rest, and was based off of the tidal volume of the subject during exercise. Consistent bag volumes were ensured using a timed switching circuit which resulted in the desired volume, given a constant rate of flow from the tank. The switching circuit and tank were calibrated for accurate volumes before each test. At the end of a normal expiration (EELV, end-expiratory lung volume) subjects were switched into the rebreathing bag and breathed the test gas for eight to ten breaths. Following each maneuver, the rebreathing bag was completely emptied via vacuum and refilled immediately prior to the next maneuver.

Custom software was used to calculate the rate of disappearance of the gases with each breath calculated from the slope of the exponential disappearance for each gas with respect to helium (31). Membrane conductance and binding of carbon monoxide to hemoglobin contribute to the diffusion capacity of the lungs for carbon monoxide (33). Unlike DLCO, DLNO is based primarily on membrane conductance (DMNO) as nitric oxide is scavenged 280 times faster by hemoglobin than carbon monoxide, causing the uptake of NO to be instantaneous ($DLNO \approx DMNO$). Therefore, DL_{NO} is considered a direct measure of membrane conductance as the diffusion resistance of the blood is insignificant (19, 20, 26, 33). DMNO was then used to calculate the D_M for carbon monoxide (DMCO) by correcting for their different diffusion constants based on molecular weight and solubility as described previously (19, 20, 26, 33). The correction factor based on molecular weight and solubility is 1.93 (16), however this correction factor is too low to produce values of V_C that are physiologically plausible. Correction factors in used in previous work have been as high as 2.49 (24, 33, 36, 41). More recently, Ceridon et al. demonstrated that a correction factor of 2.11 is most appropriate during exercise by comparing the DLNO with the more traditional

DLCO to calculate DM and V_C using DLCO at multiple oxygen tensions (6). Because this study utilized exercise, we have utilized 2.11 as the correction factor. V_C was then calculated from the DLCO measured by subtracting the resistance to diffusion associated with alveolar-capillary barrier (DMCO). Finally, we corrected for individual differences in the rate of gas uptake and binding to hemoglobin due to each individual's hemoglobin concentration and alveolar partial pressure of oxygen.

Assessment of hemoglobin

Hemoglobin concentration was assessed at the University of Arizona Medical Center Pathology Laboratory using a cyanide-free hemoglobin method on an ADVIA 2120 Hematology system. Subject's hemoglobin concentration (assessed from the baseline blood draw) was utilized in the calculation of V_C, as described above.

SCNN1A genotyping and subject grouping

The inside of each cheek of the subject was swabbed, and the swabs were immediately placed in a stabilizing buffer for storage purposes. SCNN1A was genotyped and the αENaC amino acid at position 663 encoded by each SCNN1A allele was determined at the University of Arizona Genetics Core Laboratory using a Taqman SNP assay for rs#2228576. Briefly, initial DNA quantitation and quality control was performed using PicoGreen (Life Technologies). Pre-validated primers and probe sets for TaqMan Allelic Discrimination Assay were obtained from Life Technologies. Reactions were run at 10μL, containing TaqMan Universal PCR Master Mix, No AmpEraseR UNG (Life Technologies), 10ng total DNA, and 1X Assay Mix. All samples were processed and analyzed on 7900 Real-Time PCR System (Life Technologies) with cycling conditions (95°C for 10 minutes, 50 cycles of 92°C for 15 seconds and 60°C for 1 minute) and Genotyper software (SDS system, version 2.3). Subjects were then grouped according the amino acid at position 663 of αENaC. Individuals homozygous for SCNN1A alleles encoding alanine at amino acid 663 were grouped in the AA genotype, and individuals with at least one SCNN1A allele encoding threonine at amino acid 663 of αENaC were grouped into the AT/TT genotype, as only one subject was homozygous for threonine at amino acid 663.

Statistical analysis

The SPSS statistical software package (v.19; SPSS, Inc., Chicago, IL) was used for all statistical analyses. After confirming equality of variance with a Levene's Test, a one-sided independent samples t-test was used to determine significance between genotype groups at rest and peak exercise, to determine significance between genotype groups in percent change from rest to peak exercise and also to determine significance from rest to peak exercise within genotype groups. Using Bonferroni correction, an α level of 0.025 or lower was used for determining statistical significance. All data are presented as mean ± standard deviation unless stated otherwise.

Results

Eighteen individuals with CF participated in the study (Subject demographics, Table 1). Ten subjects were homozygous for alleles resulting in an alanine at amino acid 663 (AA genotype group), and eight subjects had at least one allele resulting in a threonine (AT/TT genotype group) at amino acid 663. Only one subject was homozygous for threonine at amino acid 663 and that subject was included in the AT/TT group. There were no differences in age, height, or forced expiratory volume in one second over forced vital capacity (FEV₁/FVC) between genotype groups. Weight, body mass index (BMI), forced vital capacity (FVC) and forced expiratory volume in one second (FEV₁) were significantly lower in the AT/TT genotype group than in the AA genotype group (Table 1).

Table 2 depicts the cardiopulmonary measures taken at rest and during peak exercise. No differences were seen in cardiopulmonary parameters at rest or in response to peak exercise between the AA and AT/TT groups (Table 2). Both groups reached similar peak workloads and similar significant increases in heart rate, cardiac output, systolic blood pressure, oxygen consumption, respiratory rate, and minute ventilation in response to peak exercise were seen in both groups (Table 2).

No significant differences were seen in absolute lung diffusion measures at rest or during peak exercise; although at rest the AA group tended to have higher D_M and $DLNO/V_A$ than the AT/TT group (Table 3). The AA group demonstrated an attenuated percent increase in $DLNO$, D_M , and $DLNO/V_A$ from rest to peak compared to the AT/TT group (% increase: $DLNO= 17.8\pm 3.5$ vs. 40.6 ± 13.4 ; $DLNO/V_A= 13.8\pm 6.8$ vs. 39.7 ± 13.0 ; $D_M= 14.6\pm 3.5$ vs. 40.7 ± 13.4 , AA vs. AT/TT, $p<0.025$ for D_M , Figure 1 panels A-C). Additionally, the AA group had a larger percent increase in V_C with exercise than the AT/TT group (% increase: 87.2 ± 7.2 vs 44.9 ± 16.2 , AA vs. AT/TT, Figure 1 panel D).

Discussion

In the present study we demonstrate that genetic variation of SCNN1A manifested by changes at amino acid 663 is associated with differences in diffusing capacity of the lung in response to peak exercise in individuals with CF. Similar to previous studies, we have demonstrated an increase in both $DLCO$ and $DLNO$ with exercise in both groups (19, 33, 38, 39). Individuals with at least one allele resulting in a threonine at amino acid 663 (AT/TT group) demonstrated a significantly greater percent increase from rest to peak exercise in $DLNO$, D_M , and $DLNO/V_A$ than individuals homozygous for alleles encoding the alanine variant. The AA group had a significantly greater percent increase from rest to peak exercise in V_C . Interestingly, individuals in the AT/TT group had a significantly lower body weight, body mass index, and baseline pulmonary function (FVC, FEV_1 , and FEF_{50} % predicted). The lower resting pulmonary function in the AT/TT group may be due to the more functional ENaC resulting in a drier lung at rest which could reduce mucus clearance and result in the lower pulmonary function seen in the AT/TT group.

During exercise, the surface area for diffusion increases due to airway and capillary recruitment, and V_C tends to drive the increase in $DLCO$ and $DLNO$ due to increases in cardiac output and ventilation. Individuals in the AT/TT group demonstrated a significantly greater percent increase from rest to peak exercise in $DLNO$, D_M , and $DLNO/V_A$, bringing the AT/TT group to similar levels of conductance as the AA group. The AT/TT group demonstrated this greater percent increase in these diffusing capacity parameters despite a significantly smaller percent increase in pulmonary capillary blood volume (V_C). Given that the groups have similar maximal voluntary ventilation and alveolar volume, it is likely that these differences reflect alterations in the alveolar-capillary membrane. This suggests that the increase in conduction may be due to increased surface liquid volume in response to exercise. Improved airway hydration, through the use of hypertonic saline, has previously been shown to increase pulmonary function (FVC and FEV_1) in individuals with CF, although no diffusing capacity data was available (12). Interestingly, recent studies have demonstrated an important relationship between $DLNO$ and CT-derived structural abnormalities in patients with CF, suggesting that this technique strongly represents lung damage in this patient population(10).

There are various pathways that influence ion and, therefore, fluid regulation during exercise in the lungs. During exercise, epinephrine levels increase by up to 1000 fold causing stimulation of the β_2 -adrenergic receptor (β_2 -AR) (9). Activation of the β_2 -AR causes the release of G_{α_s} subunit which activates adenylyl cyclase resulting in the conversion of ATP

to cyclic-adenosine monophosphate (cAMP). Cyclic AMP subsequently activates protein kinase A, which phosphorylates CFTR, causing channel activation and movement of Cl⁻ to the apical side of the airway epithelial cells and a CFTR mediated inhibition of ENaC activity (4, 7). Additionally, ENaC can be directly activated by protein kinase A during exercise. ENaC activity can also be upregulated during exercise through direct activation as a result of increases in shear stress in response to increased ventilation (11, 15, 34, 40).

During exercise, ENaC activity can be inhibited via activation of the purinergic (P2Y2) pathway by adenosine triphosphate and adenosine which are released from the epithelia in response to the mechanical stress of increased ventilation during exercise (8). The application of nucleotides to the airway lumen causes increased secretion and hydration of the airway surface liquid in both healthy and CF epithelia (5). ATP and adenosine interact with P2Y2 receptors on the apical side of the membrane causing a breakdown in phosphatidylinositol 4,5-bisphosphate (PIP2). Since PIP2 is required for protein kinase C-mediated ENaC activation, the breakdown of PIP2 results in the inhibition of ENaC. Purinergic stimulation during exercise can also increase calcium activated chloride channel activity through inositol trisphosphate (IP₃) mediated release of Ca²⁺ from the endoplasmic reticulum Ca²⁺ stores (8, 22). This is important because even small changes in apical Cl⁻ and, therefore, fluid can improve pulmonary function in patients with CF. The differences seen in DLCO and DLNO from rest to peak exercise between the genetic variants at amino acid 663 of ENaC may have resulted from the response to adrenergic stimulation of CFTR resulting in ENaC inhibition or more likely through purinergic inhibition of ENaC and CaCC activation.

In healthy individuals exercise resulted in opposite changes in diffusing capacity; AA individuals demonstrated a greater increase in diffusing capacity in response to peak exercise, whereas in CF AT/TT individuals demonstrated a greater increase in diffusing capacity (3). We hypothesize that in healthy individuals the AA group have a greater capacity to increase ENaC activity in response to adrenergic stimulation leading to fluid clearance and a greater increase in diffusing capacity. In individuals with CF, exercise may lead to inhibition of the more active T663 ENaC resulting in an increased surface liquid volume which may provide beneficial improvements in diffusing capacity.

Conclusion

This study suggests that genetic variation of the alpha-subunit of ENaC is associated with differences in the diffusing capacity response to exercise in CF. We hypothesize that the greater positive change in lung diffusion in the AT/TT group, which brought this group to overall levels similar to the AA group, could be due to exercise-induced inhibition of ENaC activity being more influential for improving surface liquid depth in the AT/TT group who have what was shown to be a more active ENaC in cell work.

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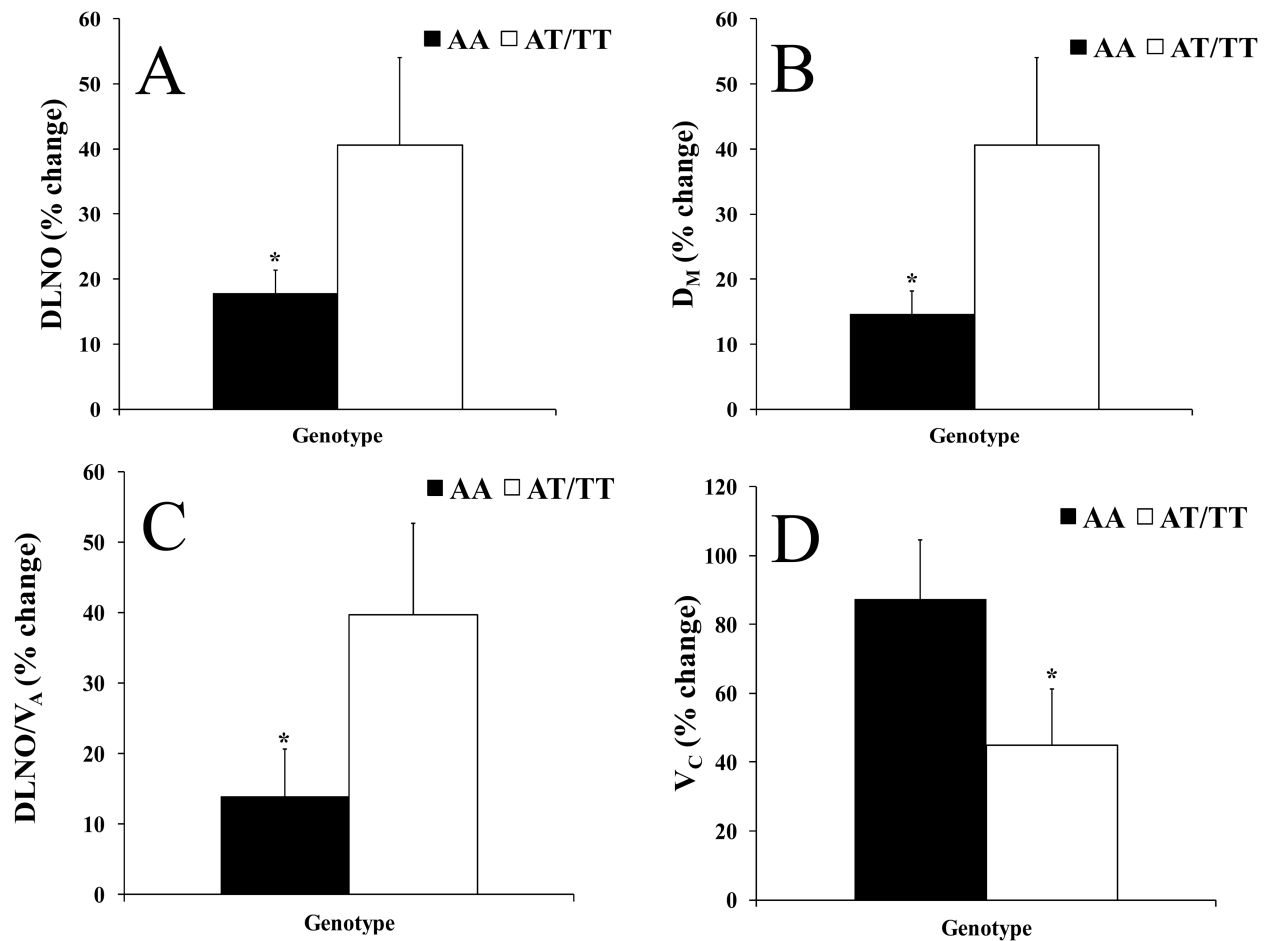


Figure 1.

Panels depict the percent increase in response to peak exercise for A) the diffusing capacity of the lung for nitric oxide (DLNO), B) alveolar-capillary membrane conductance (D_M), C) the diffusing capacity of the lung for nitric oxide corrected for alveolar volume ($DLNO/V_A$), and D) pulmonary capillary blood volume (V_C). The filled in bars represent the AA genotype (individuals homozygous for SCNN1A resulting in alanine at amino acid 663) and the open bars represent the AT/TT genotype (individuals with at least one allele resulting in threonine at amino acid 663). The error bars represent the SE of the mean. * $p < 0.05$ between genotypes, ** $p < 0.025$ between genotypes.

Table 1

Subject demographics

	AA Genotype	AT/TT Genotype
n	10	8
Age (years)	24±8	21±7
Height (cm)	168±10	169±5
Weight (kg)	70±17	56±8*
BMI (kg/m²)	25±4	20±2*
FVC (L)	3.87±1.37	3.36±1.10
FVC (%predicted)	65±37	29±29*
FEV₁ (L/sec)	2.92±1.33	2.55±.99
FEV₁ (%predicted)	60±39	25±25*
FEV₁/FVC (%)	71±15	72±8
FEF₅₀ (%predicted)	50±35	27±14
MVV (L/min)	112±48	96±28
MVV (%)	72±19	102±67

Values are mean±SD; BMI=Body mass index; FVC=forced vital capacity; FEV₁= Forced expiratory volume in 1 second; FEF₅₀=Forced expiratory flow at 50% of forced vital capacity; MVV= Maximum voluntary ventilation; AA genotype= homozygous for SCNN1A resulting in alanine at amino acid 663; AT/TT genotype= at least one copy of SCNN1A resulting in threonine at amino acid 663;

* p<0.05 between groups.

Table 2
Cardiopulmonary response to exercise

	AA Genotype		AT/TT Genotype	
	Rest	Peak Exercise	Rest	Peak Exercise
Workload (Watts)	0	100±32	0	106±44
Heart Rate	94±20	140±51 [†]	92±14	137±20 [†]
Cardiac Output (L/min)	5±2	11±4 [†]	4±1	12±4 [†]
SBP (mmHg)	107±11	144±29 [†]	106±9	137±20 [†]
DBP (mmHg)	71±9	70±16	68±8	65±14
MAP (mmHg)	83±9	94±19	81±8	89±8
VO₂ (mL/kg/min)	6±1	22±6 [†]	7±2	26±12 [†]
RER	.92±.14	1.20±.13	.89±.17	1.10±.05
RR (breaths/min)	20±5	38±8 [†]	24±7	43±11 [†]
V_T BTPS (mL)	854±352	1751±692 [†]	753±275	1498±495 [†]
V_E BTPS (L/min)	16±6	62±17 [†]	17±5	61±15 [†]

Values are mean±SD, SBP= systolic blood pressure, DBP= diastolic blood pressure, MAP= mean arterial pressure, VO₂= oxygen consumption, RER=respiratory exchange ratio, RR= respiratory rate, V_T BTPS = tidal volume standardized for temperature, pressure, and humidity, V_E BTPS= minute ventilation standardized for temperature, pressure, and humidity. AA genotype= homozygous for SCNN1A resulting in alanine at amino acid 663; AT/TT genotype= at least one copy of SCNN1A resulting in threonine at amino acid 663. No significant differences were seen between genotypes in these measures.

[†]=p<0.05 from rest to peak exercise within group.

Table 3
DLCO, DLNO, D_M , V_C , V_A , DLNO/ V_A , DLNO/DLCO, and SaO₂ in response to exercise

	AA		AT/TT	
	Rest	Peak Exercise	Rest	Peak Exercise
DLCO (mL/min/mmHg)	21± 8	27±10	17±7	24±11
DLNO (mL/min/mmHg)	70±25	82±28	53±22	74±35
D_M (mL/min/mmHg)	34±12	38±13	25±10	35±16
V_c (mL)	54±25	97±43	50±29	71±43
V_A (mL)	2215±481	2364±679	2216±734	2127±682
DLNO/V_A	31±7	35±7	25±8	33±7 [‡]
DLNO/DLCO	3.3±.3	3.0±.2	3.1±.4	3.1±.2
SaO₂ (%)	96±1	95±4	96±2	92±7

Values are mean±SD, DLCO= diffusing capacity of the lung for carbon monoxide, DLNO= diffusing capacity of the lung for nitric oxide, D_M = alveolar-capillary membrane conductance, V_C = pulmonary capillary blood volume, V_A = alveolar volume, DLNO/ V_A = diffusing capacity of the lung for nitric oxide corrected for alveolar volume, SaO₂= oxygen saturation. AA genotype= homozygous for SCNN1A resulting in alanine at amino acid 663; AT/TT genotype= at least one copy of SCNN1A resulting in threonine at amino acid 663. No significant differences were seen between genotypes.

[‡]=p<0.05 from rest to peak exercise within group.