# Assessment of the Tobacco Heating System 2.2, a candidate Modified Risk Tobacco Product, on human organotypic nasal and bronchial epithelial tissue culture using systems toxicology approach

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## INTRODUCTION

New tobacco products with the potential to reduce the risk associated with smoking are under development and require a careful safety assessment strategy.

To investigate the effect of the aerosol generated by Philip Morris's candidate Modified Risk Tobacco Product, named Tobacco Heating System 2.2 (THS2.2), organotypic tissue cultures derived from human nasal and bronchial primary epithelial cells were chosen as the test model. Tissue cultures were exposed for 28 min, at the air liquid interface, to air (sham control), THS2.2 aerosol or mainstream smoke derived from 3R4F reference cigarettes (obtained from the University of Kentucky). 3R4F smoke and THS2.2 aerosol were diluted with air to match the same nicotine level (Fig. 1). Various endpoints were collected at different time points after exposure to identify and compare the dose- and time-dependent effect of each exposure condition (Fig. 2A).

By using systems toxicology-based risk assessment approaches combining computable biological network models and gene expression changes<sup>1,2</sup>, we compared the molecular perturbations in both 3R4F smoke and THS2.2 aerosol exposure conditions. The overall assessment results reported here highlight a reduced toxicity of THS2.2 aerosol acute exposure on both nasal and bronchial epithelial tissue culture when compared to combustible cigarette smoke.



system.

Figure 2: (A) Exposure and Sample Collection.

Nasal organotypic tissue cultures derived from human primary nasal epithelial cells donor: 30 years-old male non smoker - supplier: Epithelix. **Bronchial organotypic tissue cultures** derived from human primary bronchial epithelial cells – donor 1: 23 years-old male non smoker (for gene expression analysis and inflammatory markers measurement) – supplier: MatTek, and donor 2: 28 years-old male non smoker (for histology and gene expression analysis) – supplier: Epithelix. Organotypic epithelial tissues were directly exposed at the air liquid interface for 28 min to diluted smoke from 3R4F reference cigarettes or to 60% humidified air (air-exposed controls) or to diluted THS2.2 aerosol. After exposure, inserts were incubated with fresh culture medium for 4h, 24h, 48h or 72h before measuring various endpoints. TEER: Trans Epithelial Electrical Resistance. AK: Adenylate kinase. Nicotine levels were measured in the aerosol at different dilutions (%: vol/vol with air) of 3R4F smoke and THS2.2 aerosol using the Extrelute method. Bars are means  $\pm$  SEM (N = 2-15).

Activity of cytochrome P450 (CYP) 1A1/1B1 is increased in 3R4F smoke (0.15 mg/L nicotine)-exposed nasal tissue at 48h and 72h post exposure time points. Inserts exposed to THS2.2 aerosol do not show any substantial increase of CYP activity with respect to the air control group.



Figure 5: Activity of CYP1A1/CYP1B1 following exposure in nasal organotypic tissue cultures. \* and # indicates a significant difference as compared with the corresponding air controls or 3R4F smoke concentration respectively (p-values < 0.05). The x-axis reports exposure conditions and concentration in mg/L; y-axis indicates the normalized CYP activity to positive control (TCDD) considered as 100%. N = 9-15.

A dose- and a time-dependent gene expression change is observed in 3R4F smoke-exposed <u>nasal</u> and bronchial inserts as compared to air-controls. THS2.2 aerosol exposure induces an early dose-dependent response in both tissue types. 3R4F smoke exposure always induces a stronger gene expression change than THS2.2 aerosol exposure.



Figure 6: Dose- and time-dependent effect on gene expression of 3R4F smoke- and THS2.2 aerosol-exposed nasal and bronchial inserts. Volcano plots representing global differentially expressed genes (light blue dots: down-regulated, yellow dots: up-regulated, dark blue dots: below fdr p-value of 0.05). Coefficient presents log2-based fold change. fdr: false discovery rate. N = 9-15 for nasal and N = 2-7 for bronchial.





## RESULTS







#### Reduced network perturbations of THS2.2 aerosol-exposed compared to 3R4F smoke-exposed nasal tissue at comparable level of nicotine (0.15 mg/L) and at different post-exposure time points.

|                    | 24h                              | 3R4                   | <b>₹4F</b> 48h  |   |                | 72h                                 |  |  |  |  |
|--------------------|----------------------------------|-----------------------|---|---|----------------|-------------------------------------|--|--|--|--|
| (%77)              | CST<br>(45.2%)<br>               | CP <i>R</i><br>(3.4%) | 3R4F (0.15) 48h (RBIF=14%, δ=0.83)<br>1PN<br>( <sup>16.1%</sup> ) | CS7<br>(56.7%)<br>(54.8%)<br>CEV        | CPR<br>(2.4%)  | 3R4F (0.15) 72h (RBIF=11%, 8 =0.82) | CEX<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(52.3%)<br>(5 |  |  |  |
|                    | 24h                              | THS2                  | 2.2   | 48h                                     |                |                                     | 72h  |  |  |  |
| (62.7%)            | CST<br>(15.6%)<br>(%6.6k)<br>VHO | CPR<br>(1.8%)         | THS2.2 (0.15) 48h (RBIF=0%, δ=-0.01)<br><b>IPN</b><br>(0%)        | CST<br>(47%)<br>(47%)<br>(%1.6%)<br>CEV | CPR<br>(11.4%) | THS2.2 (0.15) 72h (RBIF=0%, 8 =0.3) | CST<br>(78.4%)<br>(%0)<br>(%0)<br>(%0)<br>(%0)<br>(%0)<br>(%0)<br>(%0)<br>(%   |  |  |  |
| onary Inflammation |                                  |                       | Cell Proliferation  |   |                |                                     | Cell Stress  |  |  |  |

Figure 7: Quantification of the perturbations of different biological networks using gene fold-changes as an input and the NPA method<sup>3</sup> computational approach. The surface area of each slice is proportional to the contribution within a particular treatment of each network perturbation (shown in percentage). The pie chart represents the distribution of the sum of the contributions o each network across all treatment groups. N = 9-15.

- time and dose (Figure 5).
- pulmonary inflammation (IPN) (Figure 7).
- cultures (VEGFA, TIMP1, siCAM1, MMP1).

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## The release of various inflammatory mediators in the <u>nasal</u> tissue cultures was significantly increased

| 48h         | 72h  | 24h            | 48h | 72h | 24h  | 48h     | 72h |          |
|-------------|------|----------------|-----|-----|------|---------|-----|----------|
|             |      |                |     |     |      |         |     | -        |
| 0.9         | 0.7  | 1.4            | 1.1 | 1.1 | 2.1  | 1.8     | 1.5 | - VEGFA  |
| 1.2         | 1.1  | 1.0            | 1.2 | 0.8 | 0.9  | 0.8     | 0.9 | - TNFA   |
| 0.9         | 1.3  | 1.2            | 1.1 | 1.2 | 1.3  | 1.0     | 1.8 | - TIMP1  |
| 1.1         | 0.9  | 1.1            | 1.1 | 0.8 | 1.3  | 0.9     | 0.9 | - sICAM1 |
| 1.2         | 1.0  | 1.2            | 1.0 | 0.9 | 1.2  | 0.9     | 1.0 | – MMP-9  |
| 1.1         | 0.8  | 1.1            | 1.2 | 0.8 | 1.2  | 1.2     | 1.1 | – MMP-1  |
| 1.0         | 0.8  | 0.9            | 0.9 | 0.6 | 1.1  | 0.9     | 1.1 | - CCL20  |
| 1.2         | 1.0  | 1.2            | 1.3 | 0.9 | 1.4  | 1.2     | 1.0 | - CXCL8  |
| 1.3         | 1.1  | 1.3            | 1.4 | 0.7 | 1.2  | 0.8     | 0.8 | – IL6    |
| 1.1         | 0.8  | 1.0            | 1.0 | 0.7 | 1.0  | 1.2     | 1.0 | - CXCL1P |
| 1.0         | 1.2  | 0.9            | 1.2 | 0.9 | 0.7  | 1.1     | 1.0 | – EGF    |
| 52.2 (0     | .15) | THS2.2 (0.25)  |     |     | THS  | 2,2 (0  | -   |          |
| 2.2 (0.10)  |      | 11132.2 (0.23) |     |     |      | (0      |     |          |
| HS2.2 (Air) |      | – THS2.2 (Air) |     |     | — T⊦ | IS2.2 ( |     |          |
|             | . ,  |                |     |     |      |         | . , |          |

Figure 8: Multi-analyte profiling (MAP) of proinflammatory mediators. Fold-changes of various mediators (listed in the y-axis, right) were measured in the basolateral media of tissue cultures by Luminex technology; x-axis indicates the ratio between exposure condition/air control. Dark and light grey colors indicate the significant increase and decrease, respectively, between the contrasts (p-values < 0.05). N = 9-15.



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| have lower proportion of the FoxJ1 positive stained cells                    |            |         |   |                          |             |                                |          |          |                           |   |  |
|--|------------|---------|---|--------------------------|-------------|--------------------------------|----------|----------|---------------------------|---|--|
| 72h after exposure whereas THS2.2 aerosol-exposed                            |            |         |   |                          |             |                                |          |          |                           |   |  |
| tissues had lower proportion of the Ki67-positive stained                    |            |         |   |                          |             |                                |          |          |                           |   |  |
|  |            |         |   |                          |             |                                |          |          |                           |   |  |
| cells 48 and 72h after exposure.   |            |         |   |                          |             |                                |          |          |                           |   |  |
| Δ  |            |         |   |                          |             |                                |          |          |                           |   |  |
| A  |            |         | Air                                       | Contro                   |             | 3R4F (0.15)                    |          |          |                           |   |  |
| H&E/ Alcian<br>Blue  |            |         | the Marson                                | 10000                    | Sileal L    | ator of sugar and sugar boards |          |          |                           |   |  |
|  |            |         | 644 2040                                  |                          | Distance in |                                |          |          |                           |   |  |
|  |            | i i     | 20 40 60 80                               | 100 µm                   | j j         | ο 20 40 60 80 100μm            |          |          |                           |   |  |
|  | F          | oxi1    | 9- 8- 8 - 8 - 8                           | 200000                   |             | 186 319/10                     | Van 95'. | Blacker  | 80                        |   |  |
| (  | ciliated   | cells)  | 20 00 00                                  |                          |             |                                |          |          |                           |   |  |
|  |            | 6       | 20 40 60 8                                | 0 100 µm                 | i           | <u>0 20 40 60 80 100 µm</u>    |          |          |                           |   |  |
|  | ĸ          | (i-67 闎 | 1. J. 11010.0.                            |                          | Starting by |                                |          |          |                           |   |  |
|  | (prolifer) | ating 📗 |   |                          | 1           |                                |          |          |                           |   |  |
|  |            | , i     | 20 40 60 80 100 µ                         | n                        | 5           |                                |          |          |                           |   |  |
|  |            |         |   |                          | in when a   | ma walde Sign                  |          |          |                           |   |  |
|  | (basal d   | P63     | 6101 100 10 10 10 10 10 10 10 10 10 10 10 | Contraction of the state | 41.00       |                                |          |          |                           |   |  |
| _  | (          | 5       | 20 40 60 80 100 µm                        |                          |             | 0 20 40 60 80 100µm            |          |          |                           |   |  |
| В  |            | 1.1     | 0.8                                       | 0.9                      | 0.9         | FoxJ1                          | ſ        | 13       | <b>T</b> EON LI           | 1 |  |
|  | 48h        | 0.7     | 0.7                                       | 0.7                      | 0.6         | Ki67                           | 48h      | 1.0      | – Ki67                    | L |  |
|  |            | 1.1     | 1.0                                       | 1.0                      | 1.0         | P63                            |          | 1.0      | P63                       |   |  |
|  | 72h        | 0.6     | 1.0                                       | 1.1                      | 0.9         | FoxJ1                          |          | 0.6      | FoxJ                      | t |  |
|  |            | 0.9     | 0.9                                       | 0.7                      | 1.0         | $-\frac{K107}{P63}$            | 72h      | 1.0      | $-\frac{K_{167}}{P_{63}}$ |   |  |
|  | I          | 3D/F    | THS2 2                                    | THS2 2                   | тнс2 2      | <b>,</b>                       | 3        | P/F (0.1 | <b>_ 1</b> 05             |   |  |
|  |            | (0.15)  | (0.15)                                    | (0.25)                   | (0.44)      | -                              | T        | HS2.2 (0 | .15)                      |   |  |
| Figure 4: (A) Representative Microscopy images and (B)                       |            |         |   |                          |             |                                |          |          |                           |   |  |
| quantitative analysis of FoxJ1, Ki67, and P63 immunostaining                 |            |         |   |                          |             |                                |          |          |                           |   |  |
| after 48h and 72h after exposure. Light grey colors in the tables            |            |         |   |                          |             |                                |          |          |                           |   |  |
| indicate the significant decreased contrasts (p-values $< 0.05$ ). N = 9-15. |            |         |   |                          |             |                                |          |          |                           |   |  |

3R4F smoke (0.15 mg/L nicotine)-exposed nasal tissues

### CONCLUSIONS

Human nasal and bronchial tissue cultures were successfully exposed for 28 min to air, to diluted 3R4F cigarette smoke and THS2.2 aerosol at similar nicotine concentration (Figure 2A).

The exposure to 3R4F smoke induced nasal tissue damage. On the contrary, THS2.2 aerosol exposure did not result in any damage at comparable nicotine concentration (Figure 3A, B, C).

Cell proliferation (Ki-67) is reduced in THS2.2 aerosol-exposed nasal tissue cultures, whereas the proportion of ciliated cells (Foxj1) is reduced in 3R4F smoke-exposed cultures after 72h from exposure (Figure 4).

The activity of CYP1A1/1B1 is not substantially increased by THS2.2 aerosol exposure in nasal epithelial cultures, whereas exposure to 3R4F smoke causes a significant increase, depending on the post –exposure

A dose- and a time-dependent effect was observed after 3R4F smoke exposure at the gene expression level; gene expression changes appear mainly as an early response (4h post-exposure time point) after THS2.2 aerosol exposure in both nasal and bronchial exposed inserts; lower gene expression changes were observed for THS2.2 aerosol when compared to 3R4F smoke at all the post exposure time point (Figure 6).

The networks mostly perturbed by 3R4F smoke exposure were the cell stress (CST), cell fate (CFA) and the

The release of various inflammatory mediators (VEGFA, TNFA, TIMP1, siCAM1, MMP-9, MMP-1, CCL20, CXCL8, IL6, EGF) was increased after 3R4F smoke exposure compared to air in nasal tissue cultures in a time- and dose- dependent way. Less analytes were found upregulated in THS2.2 aerosol exposed tissue

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