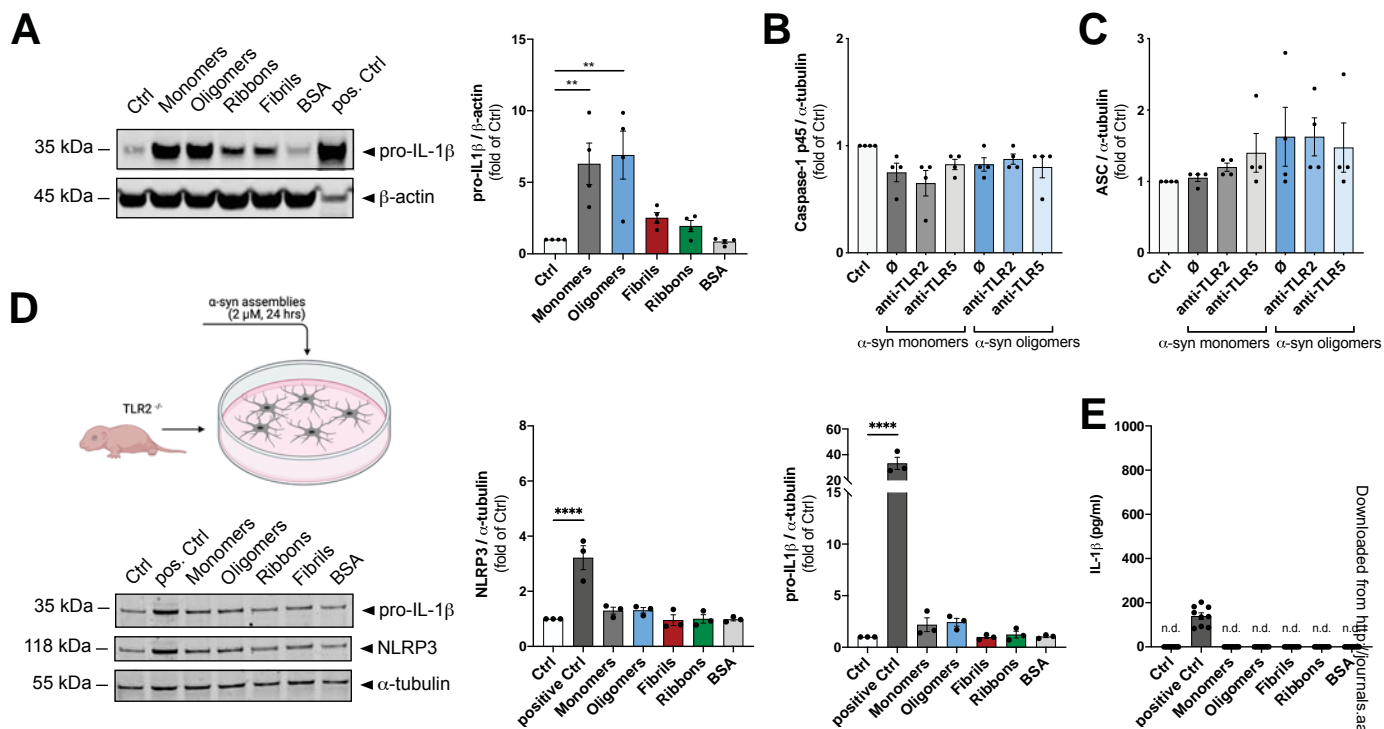


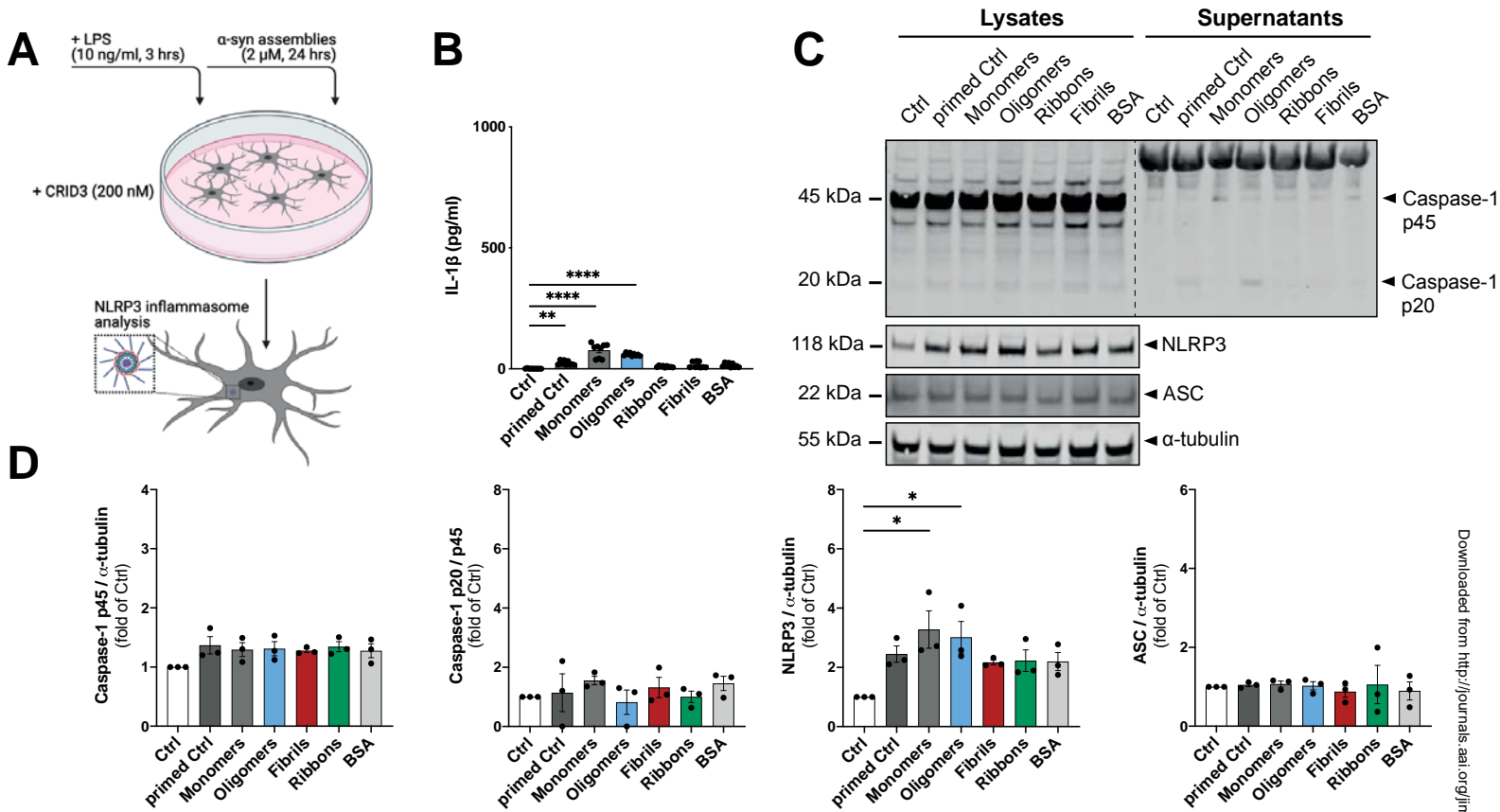
**Figure S1**

(A) Schematic drawing of the experimental setup used in this study. (B) Quantification of the different  $\alpha$ -syn forms for endotoxin-free preparations as endotoxin-free. ( $n=3$  independent experiments with triplicated measurements). (C) Quantification of IL-1 $\beta$  levels in conditioned medium of primary microglia treated for 24 hrs with different concentrations of distinct  $\alpha$ -syn forms. ( $n=4$  independent experiments with duplicated measurements). Quantification of the relative metabolic activity (D) and the relative LDH release (E) by microglia treated with 2  $\mu$ M  $\alpha$ -syn under basal conditions or after pre-exposure to LPS. See also Fig. 1. All graphs are presented as mean  $\pm$  SEM and were analysed by one-way ANOVA followed by Tukey's multiple comparison post hoc test. Levels of significance are: \*\*\*\* $p < 0.0001$ , \*\*\* $p < 0.001$ , \*\* $p < 0.01$ , \* $p < 0.05$ .



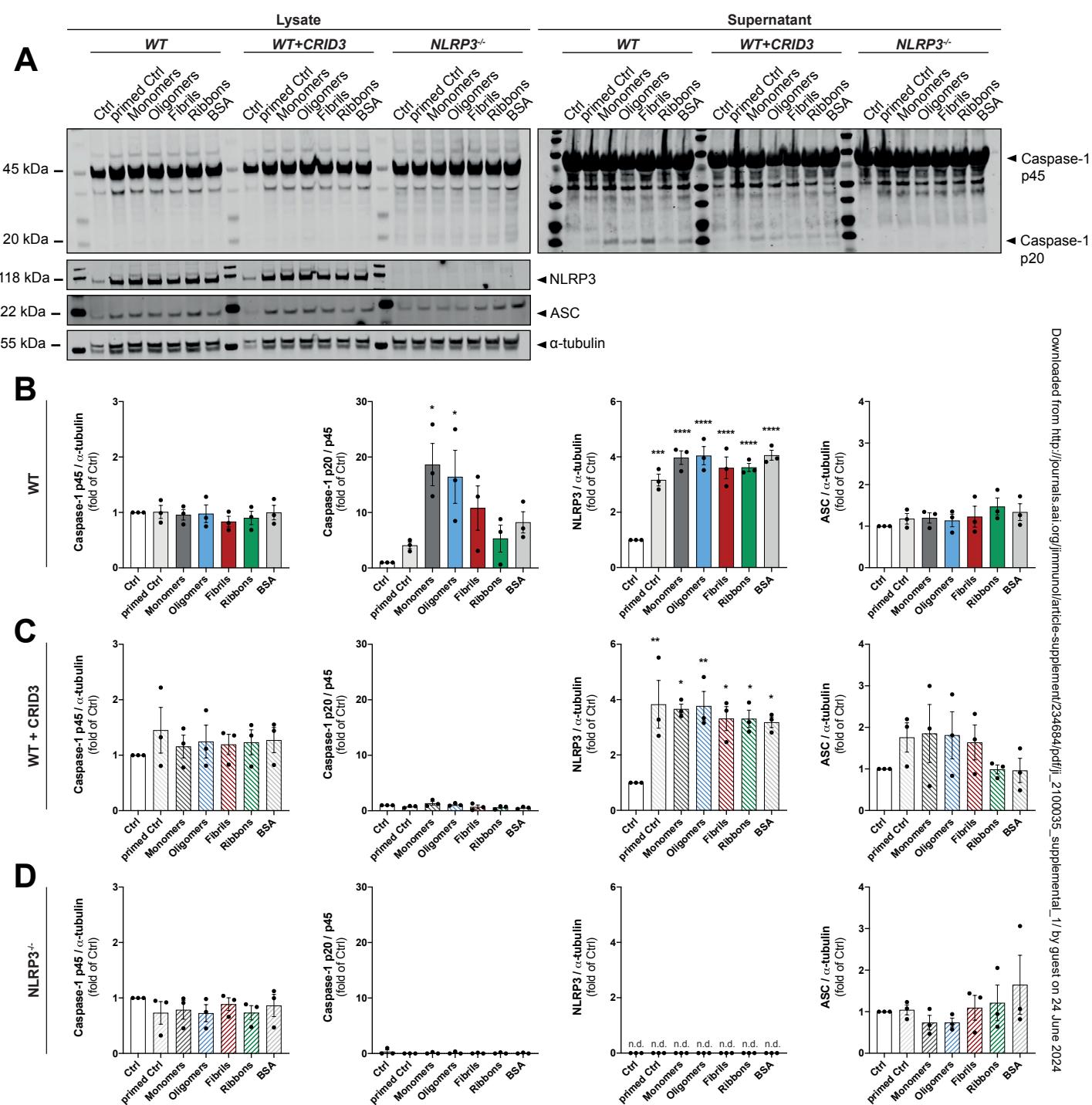
**Figure S2**

**(A)** Immunoblot analysis and quantifications of pro-IL1β levels in microglia treated with different α-syn assemblies. Quantification of caspase-1 p45 **(B)** and ASC **(C)** expression upon treatment with TLR2 and TLR5 neutralizing antibodies. (n=4 independent experiments). **(D)** Immunoblot analysis and quantifications of pro-IL1β and NLRP3 levels in microglia derived from TLR2<sup>-/-</sup> mice treated with different α-syn assemblies. (n=3 independent experiments). **(E)** Quantification of IL-1β levels in conditioned medium of primary microglia derived from TLR2<sup>-/-</sup> mice treated for 24 hrs with 2 μM distinct α-syn forms and 200 nM CRID3. (n=3 independent experiments with triplicated measurements). See also Fig. 2. All graphs are presented as mean ± SEM and were analysed by one-way ANOVA followed by Tukey's multiple comparison post hoc test **(A, D-E)** or Kruskal-Wallis test for nonparametric data **(B-C)**. Levels of significance are: \*\*\*\*p < 0.0001, \*\*\*p < 0.001, \*\*p < 0.01, \*p < 0.05.



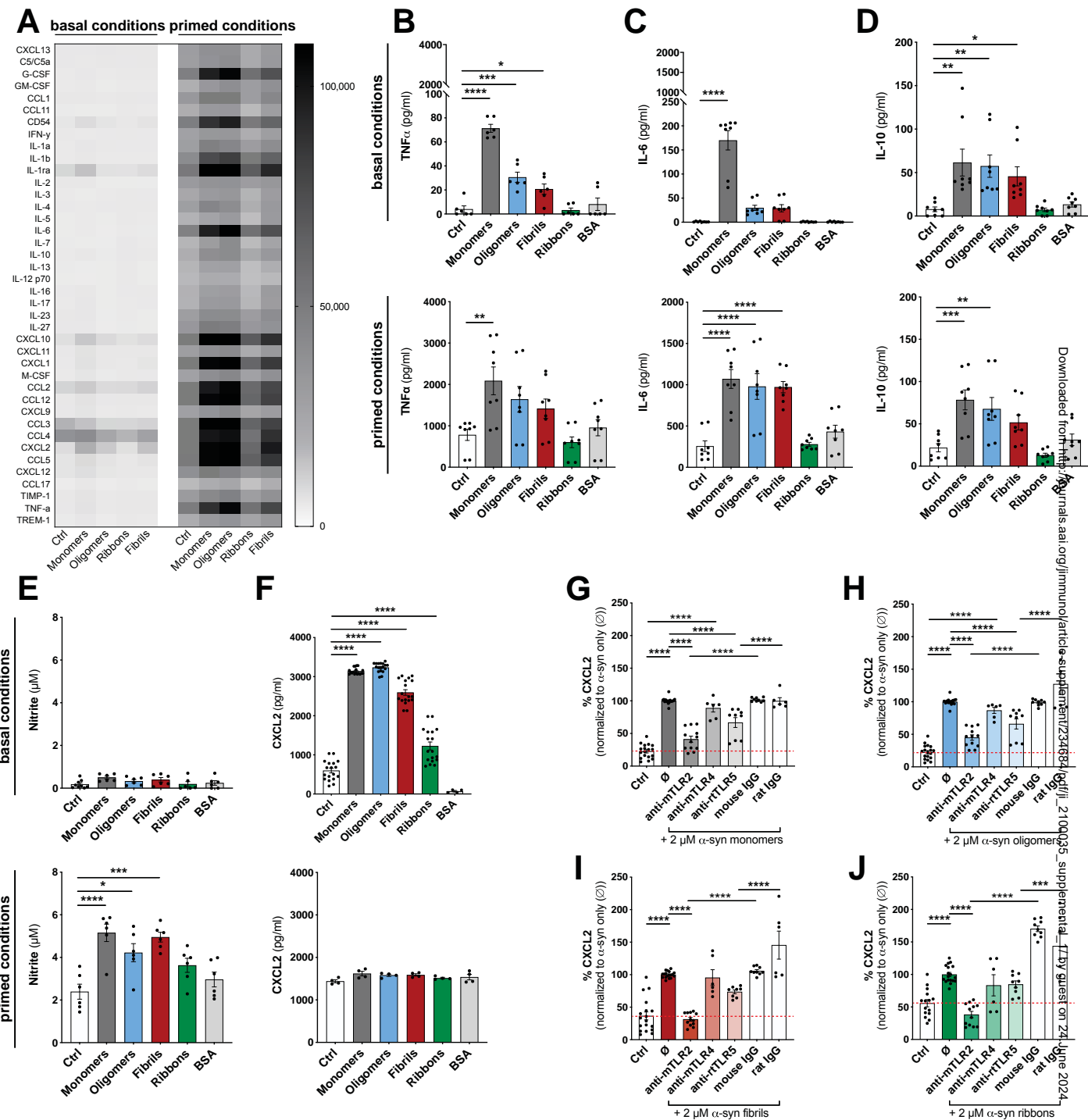
**Figure S3**

**(A)** Schematic drawing of the experimental setup used for the following experiments **(B-D)**. **(B)** Quantification of IL-1β levels in conditioned medium of LPS-primed primary microglia treated for 24 hrs with 2 μM distinct α-syn forms and 200 nM CRID3. (n=3 independent experiments with triplicated measurements). **(C-D)** Immunoblot and quantification of primed microglia cell lysates and supernatants exposed to 2 μM α-syn and 200 nM CRID3 for 24 hrs, stained for caspase-1, NLRP3, ASC and α-tubulin. All graphs are presented as mean ± SEM and were analysed by one-way ANOVA followed by Tukey's multiple comparison post hoc test. Levels of significance are: \*\*\*\*p < 0.0001, \*\*p < 0.01, \*p < 0.05.



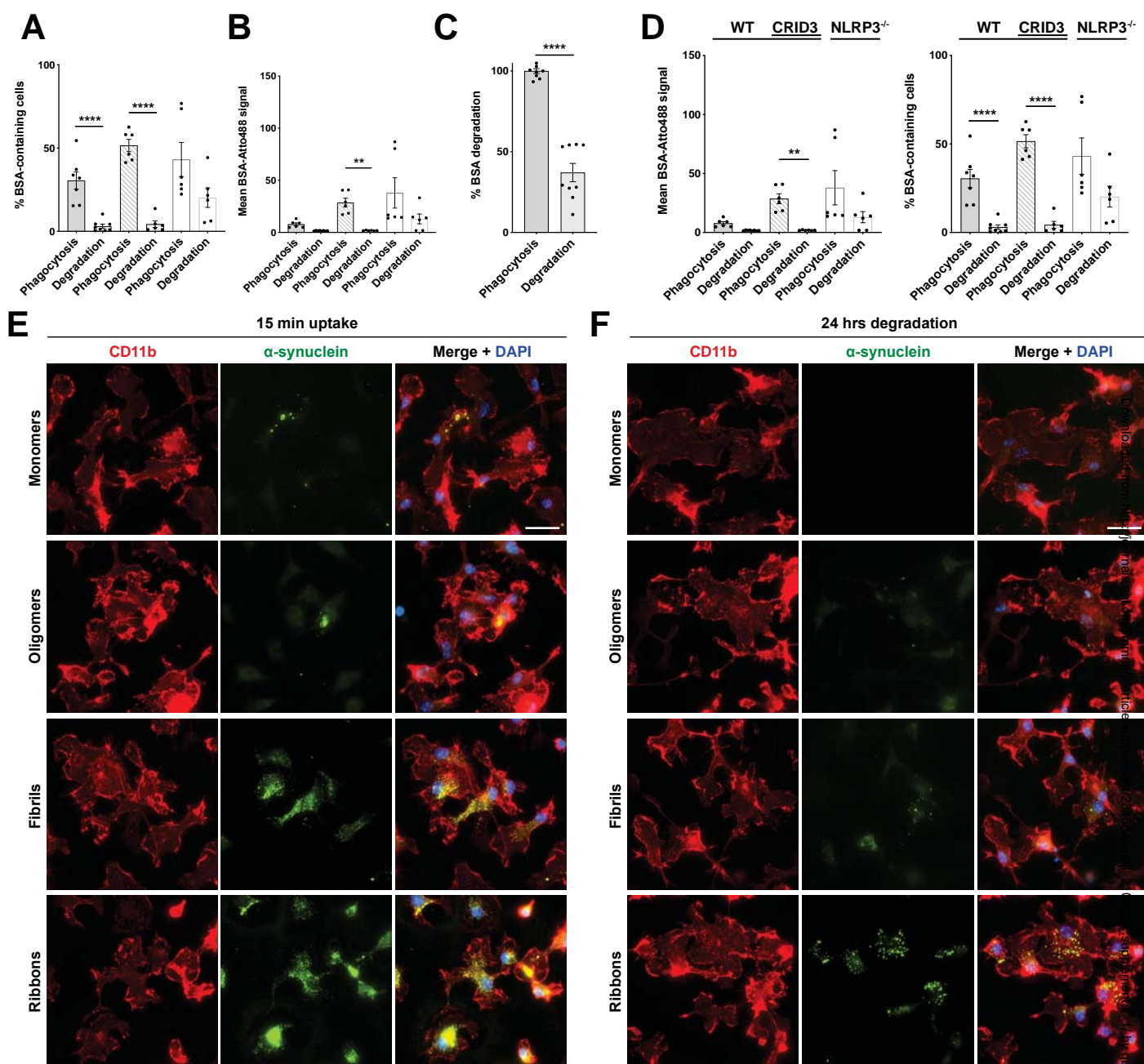
**Figure S4**

(A) Immunoblot of cell lysates (left) and supernatants (right) of primary microglia derived from WT or NLRP3<sup>-/-</sup> mice exposed to 2  $\mu$ M  $\alpha$ -syn and 200 nM CRID3 for 24 hrs. Blots were stained for caspase-1, NLRP3, ASC and  $\alpha$ -tubulin. Quantification of Caspase-1 p45, cleaved Caspase-1 p20, NLRP3, and ASC in WT (B), WT+CRID3 (C), and NLRP3<sup>-/-</sup> (D). n=3 independent experiments. All graphs are presented as mean  $\pm$  SEM and were analysed by one-way ANOVA followed by Tukey's multiple comparison post hoc test. Levels of significance are: \*\*\*\*p < 0.0001, \*\*\*p < 0.001, \*\*p < 0.01, \*p < 0.05.



**Figure S5**

(A) Proteome profiling array on supernatants of primary microglia treated with 2  $\mu\text{M}$  distinct  $\alpha\text{-syn}$  forms without or with pre-exposure to LPS for 3 hrs. Quantification of TNF $\alpha$  (B), IL-6 (C), IL-10 (D), Nitrite (E), and CXCL2 (F) in supernatants of primary microglia treated with 2  $\mu\text{M}$  distinct  $\alpha\text{-syn}$  forms without (upper panel) or with (lower panel) pre-exposure to LPS for 3 hrs. (n=3-4 independent experiments with single, duplicate or triplicate measurements). CXCL2 levels in conditioned medium of unprimed primary microglia treated for 24 hrs with TLR2-, TLR4, and TLR5-neutralizing antibodies or the respective isotype controls in parallel to stimulation with  $\alpha\text{-syn}$  monomers (G), oligomers (H), fibrils (I) and ribbons (J). (n=4-6 independent experiments with duplicate or triplicate measurements). All graphs are presented as mean  $\pm$  SEM and were analysed by one-way ANOVA followed by Tukey's multiple comparison post hoc test. Levels of significance are: \*\*\*\*p < 0.0001, \*\*\*p < 0.001, \*\*p < 0.01, \*p < 0.05.



**Figure S6**

Quantification of the percent of phagocytic cells (**A**), the phagocytic index (**B**), and the percentage of degradation (**C**) of primary microglia exposed to BSA as protein control. (n=5-6 independent experiments). For comparison see also Fig. 4 and 5. (**D**) Quantification and comparison of phagocytosis and degradation of BSA by WT microglia (filled bars), WT microglia treated with the NLRP3 inflammasome inhibitor CRID3 (striped bars), or NLRP3-deficient microglia (empty bars). Representative immunostainings for CD11b-positive microglia (red) and distinct forms of  $\alpha$ -syn-Atto488 (green) after 15 minutes of phagocytosis (**E**) or 24 hours of degradation (**F**). All graphs are presented as mean  $\pm$  SEM and were analysed by one-way ANOVA followed by Tukey's multiple comparison post hoc test. Levels of significance are: \*\*\*\*p < 0.0001, \*\*\*p < 0.001, \*\*p < 0.01, \*p < 0.05. Scale bar: 20  $\mu$ m.

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