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Supplement of

Global change in streamflow extremes under climate change over the 21st century

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Supplementary Information

The average value of change in streamflow extremes for each GCM-GHM combination, under RCP8.5 and RCP2.6 warming scenarios, are presented individually in the Tables S2 and S3, respectively. It should be noted that the results presented in the Tables S2 and S3, if averaged over the models for each quadrant, may result in different values from the multi-model average values reported in the main text. For the values reported in the main text, the results are first averaged over the models for each grid cell to create a multi-model global matrix. The grid-cells are then averaged over each quadrant, weighted by grid cell area. Assignment of each grid cell to the specified quadrants is based on the multi-model averages. For any single model datasets, grid cells (especially the ones with small multimodel mean projected changes) may fall in different quadrants, due to the disagreement among the GCMs and GHMs on projection of changes.

Some ISI-MIP hydrological impact models do not simulate streamflow for some grid cells for which other models do. This is due to the differences in the utilized global river network. Hence, a minority of grid cells do not have data from all hydrological models. However, grid cells with less than 20 (out of 25) available models are excluded from the calculations.

The results presented in the main text are based on the 95th and 5th percentiles of streamflow. Maps of changes in the median of streamflow are also presented in this supplement, along with the maps of changes in 95th and 5th percentiles.

The two-sample t-test (Snedecor and Cochran, 1989) is used in this study to quantify the statistical significance level of difference between the means of the 20C and 21C streamflow time series:

$$T = \frac{\bar{Q}_{21C} - \bar{Q}_{20C}}{\sqrt{\frac{\sigma_{21C}}{N_{21C}} + \frac{\sigma_{20C}}{N_{20C}}}} \quad \text{Eq.S1}$$

where \bar{Q}_{21C} and \bar{Q}_{20C} are the mean (high or low) streamflow, σ_{21C} and σ_{20C} are the streamflow time series variances, and N_{21C} and N_{20C} are the streamflow time series sample sizes (equal to 30-years in this study), in 21C and 20C, respectively. The means are significantly different if the absolute value of T is larger than the critical value of the t-distribution with ν degrees of freedom, with ν in an equal variance assumption being equal to $N_{21C} + N_{20C} - 2$. The value of Q here can be P5 or P95, meaning that we have calculated the T based on the P5 time series in 20C and 21C, and similarly based on the P95 time series. The critical value is defined based on the confidence level of significance and sample size, and at 95% level for a sample size of 30 it is equal to 2.042. The percentage of land area with statistically significant change (at 95% confidence level) is reported in this study.

Supplementary References:

Schewe, J., Heinke, J., Gerten, D., Haddeland, I., Arnell, N. W., Clark, D. B., Dankers, R., Eisner, S., Fekete, B. M., Colón-González, F. J., Gosling, S. N., Kim, H., Liu, X., Masaki, Y., Portmann, F. T., Satoh, Y., Stacke, T., Tang, Q., Wada, Y., Wisser, D., Albrecht, T., Frieler, K., Piontek, F., Warszawski, L. and Kabat, P.: Multimodel assessment of water scarcity under climate change, *Proc. Natl. Acad. Sci.*, 111(9), 3245–3250, doi:10.1073/pnas.1222460110, 2013.

Snedecor, G. W. and Cochran, W. G.: *Statistical methods*, 8thEdn, Iowa State University Press, Iowa., 1989.

Table S1. Main characteristics of the global hydrological models used in this study (obtained from Schewe et al., 2013). LW: downwelling long-wave radiation; LWn: net long-wave radiation; P: precipitation rate (rain and snow calculated in the model); Q: air specific humidity; R: rainfall rate; RH: relative humidity; S: snowfall rate; SP: surface pressure; SW: downwelling shortwave radiation; T: air temperature; Tmax(min): daily maximum (minimum) air temperature; W: wind speed.

| Model name | Time step | Meteorological forcing | Energy balance | Evaporation scheme | Runoff scheme | Snow scheme | Vegetation dynamics | CO2 effect |
|-------------|-----------|-------------------------|----------------|--------------------|---------------------------------|---|---------------------|------------|
| DBH | 1hour | P, T, W, Q, LW, SW, SP | Yes | Energy balance | Infiltration excess | Energy balance | No | Constant |
| LPJmL | Daily | P, T, LWn, SW | Yes | Priestley-Taylor | Saturation excess | Degree-day | Yes | Varying |
| Mac-PDM | Daily | P, T, W, Q, LWn, SW, SP | No | Penman-Monteith | Saturation excess, non-linear | Degree-day | No | No |
| PCR-GLO BWB | Daily | P, T | No | Hamon | Saturation Excess Beta Function | Degree-day | No | No |
| WBM | Daily | P, T | No | Hamon | Saturation Excess | Empirical temp and Precip-based formula | No | No |

Table S2. Normalized change in high and low streamflow extremes, averaged for each quadrant. Results presented for each model under RCP8.5 scenario. The results can also be reverted to the relative change in percentage for more distinct comparison (Figure S1).

| RCP8.5 | | Quad. 1. increased high extreme and decreased low extreme | | Quad. 2. increased high and low extreme | | Quad. 3. decreased high and low extreme | | Quad. 4. decreased high extreme and increased low extreme | |
|------------------------|------------|---|--------------------|---|--------------------|---|--------------------|---|--------------------|
| GCM | GHM | Change in high ext. | Change in low ext. | Change in high ext. | Change in low ext. | Change in high ext. | Change in low ext. | Change in high ext. | Change in low ext. |
| GFDL-ESM2 _m | WBM | 0.072 | 0.178 | 0.196 | -0.291 | -0.285 | 0.451 | -0.105 | -0.171 |
| | MacPDM | 0.056 | 0.108 | 0.152 | -0.206 | -0.167 | 0.294 | -0.066 | -0.189 |
| | PCR-GLOBWB | 0.076 | 0.175 | 0.150 | -0.186 | -0.235 | 0.393 | -0.121 | -0.172 |
| | DBH | 0.079 | 0.074 | 0.129 | -0.214 | -0.161 | 0.169 | -0.101 | -0.314 |
| | LPJmL | 0.079 | 0.191 | 0.129 | -0.231 | -0.161 | 0.405 | -0.088 | -0.251 |
| HadGEM2-ES | WBM | 0.091 | 0.287 | 0.306 | -0.477 | -0.271 | 0.496 | -0.095 | -0.246 |
| | MacPDM | 0.051 | 0.077 | 0.172 | -0.224 | -0.148 | 0.270 | -0.080 | -0.192 |
| | PCR-GLOBWB | 0.102 | 0.250 | 0.226 | -0.292 | -0.268 | 0.474 | -0.142 | -0.218 |
| | DBH | 0.103 | 0.087 | 0.166 | -0.301 | -0.153 | 0.203 | -0.094 | -0.415 |
| | LPJmL | 0.110 | 0.220 | 0.176 | -0.267 | -0.141 | 0.412 | -0.096 | -0.288 |
| IPSL-CM5A-LR | WBM | 0.095 | 0.226 | 0.278 | -0.423 | -0.357 | 0.563 | -0.114 | -0.201 |
| | MacPDM | 0.058 | 0.087 | 0.179 | -0.216 | -0.190 | 0.302 | -0.066 | -0.143 |
| | PCR-GLOBWB | 0.108 | 0.241 | 0.241 | -0.254 | -0.289 | 0.481 | -0.155 | -0.220 |
| | DBH | 0.112 | 0.081 | 0.200 | -0.281 | -0.191 | 0.200 | -0.126 | -0.444 |
| | LPJmL | 0.128 | 0.211 | 0.188 | -0.278 | -0.187 | 0.394 | -0.110 | -0.279 |
| MIROC-ESM | WBM | 0.092 | 0.215 | 0.267 | -0.423 | -0.340 | 0.527 | -0.089 | -0.204 |
| | MacPDM | 0.041 | 0.061 | 0.116 | -0.134 | -0.150 | 0.259 | -0.038 | -0.087 |
| | PCR-GLOBWB | 0.103 | 0.261 | 0.223 | -0.270 | -0.272 | 0.443 | -0.137 | -0.194 |
| | DBH | 0.101 | 0.089 | 0.173 | -0.322 | -0.180 | 0.182 | -0.109 | -0.397 |
| | LPJmL | 0.116 | 0.200 | 0.160 | -0.299 | -0.164 | 0.389 | -0.104 | -0.299 |
| NorESM1-M | WBM | 0.081 | 0.215 | 0.189 | -0.345 | -0.255 | 0.429 | -0.082 | -0.204 |
| | MacPDM | 0.038 | 0.063 | 0.111 | -0.151 | -0.127 | 0.218 | -0.077 | -0.111 |
| | PCR-GLOBWB | 0.072 | 0.210 | 0.152 | -0.218 | -0.219 | 0.377 | -0.105 | -0.191 |
| | DBH | 0.074 | 0.058 | 0.121 | -0.202 | -0.140 | 0.122 | -0.072 | -0.356 |
| | LPJmL | 0.077 | 0.201 | 0.120 | -0.237 | -0.133 | 0.309 | -0.103 | -0.244 |

Table S3. Normalized change in high and low streamflow extremes, averaged for each quadrant. Results presented for each model under RCP2.6 scenario. The results can also be reverted to the relative change in percentage for more distinct comparison (Figure S1).

| RCP2.6 | | Quad. 1. increased high extreme and decreased low extreme | | Quad. 2. increased high and low extreme | | Quad. 3. decreased high and low extreme | | Quad. 4. decreased high extreme and increased low extreme | |
|------------------------|------------|---|--------------------|---|--------------------|---|--------------------|---|--------------------|
| GCM | GHM | Change in high ext. | Change in low ext. | Change in high ext. | Change in low ext. | Change in high ext. | Change in low ext. | Change in high ext. | Change in low ext. |
| GFDL-ESM2 _m | WBM | 0.067 | 0.143 | 0.153 | -0.224 | -0.153 | 0.270 | -0.063 | -0.134 |
| | MacPDM | 0.035 | 0.108 | 0.094 | -0.137 | -0.104 | 0.171 | -0.032 | -0.081 |
| | PCR-GLOBWB | 0.055 | 0.120 | 0.110 | -0.173 | -0.123 | 0.228 | -0.065 | -0.125 |
| | DBH | 0.046 | 0.037 | 0.086 | -0.116 | -0.077 | 0.068 | -0.044 | -0.128 |
| | LPJmL | 0.042 | 0.162 | 0.078 | -0.158 | -0.065 | 0.237 | -0.045 | -0.178 |
| HadGEM2-ES | WBM | 0.069 | 0.149 | 0.184 | -0.297 | -0.151 | 0.279 | -0.062 | -0.141 |
| | MacPDM | 0.031 | 0.052 | 0.097 | -0.136 | -0.068 | 0.125 | -0.033 | -0.072 |
| | PCR-GLOBWB | 0.072 | 0.154 | 0.141 | -0.181 | -0.145 | 0.271 | -0.080 | -0.143 |
| | DBH | 0.059 | 0.048 | 0.105 | -0.142 | -0.068 | 0.082 | -0.044 | -0.163 |
| | LPJmL | 0.073 | 0.165 | 0.106 | -0.183 | -0.061 | 0.233 | -0.049 | -0.182 |
| IPSL-CM5A-LR | WBM | 0.055 | 0.124 | 0.141 | -0.235 | -0.170 | 0.278 | -0.088 | -0.138 |
| | MacPDM | 0.033 | 0.054 | 0.082 | -0.103 | -0.088 | 0.140 | -0.026 | -0.071 |
| | PCR-GLOBWB | 0.055 | 0.131 | 0.108 | -0.144 | -0.161 | 0.274 | -0.091 | -0.149 |
| | DBH | 0.055 | 0.046 | 0.098 | -0.142 | -0.092 | 0.095 | -0.045 | -0.190 |
| | LPJmL | 0.060 | 0.165 | 0.084 | -0.180 | -0.078 | 0.251 | -0.055 | -0.162 |
| MIROC-ESM | WBM | 0.056 | 0.156 | 0.175 | -0.263 | -0.208 | 0.329 | -0.063 | -0.135 |
| | MacPDM | 0.031 | 0.040 | 0.073 | -0.106 | -0.074 | 0.126 | -0.019 | -0.047 |
| | PCR-GLOBWB | 0.063 | 0.159 | 0.146 | -0.196 | -0.176 | 0.289 | -0.072 | -0.135 |
| | DBH | 0.057 | 0.059 | 0.114 | -0.157 | -0.102 | 0.093 | -0.074 | -0.201 |
| | LPJmL | 0.063 | 0.172 | 0.091 | -0.190 | -0.094 | 0.272 | -0.067 | -0.184 |
| NorESM1-M | WBM | 0.045 | 0.116 | 0.129 | -0.205 | -0.147 | 0.235 | -0.049 | -0.112 |
| | MacPDM | 0.023 | 0.048 | 0.064 | -0.082 | -0.063 | 0.118 | -0.024 | -0.054 |
| | PCR-GLOBWB | 0.051 | 0.124 | 0.087 | -0.129 | -0.133 | 0.239 | -0.054 | -0.112 |
| | DBH | 0.050 | 0.036 | 0.069 | -0.114 | -0.082 | 0.068 | -0.036 | -0.122 |
| | LPJmL | 0.041 | 0.155 | 0.062 | -0.145 | -0.056 | 0.211 | -0.048 | -0.149 |

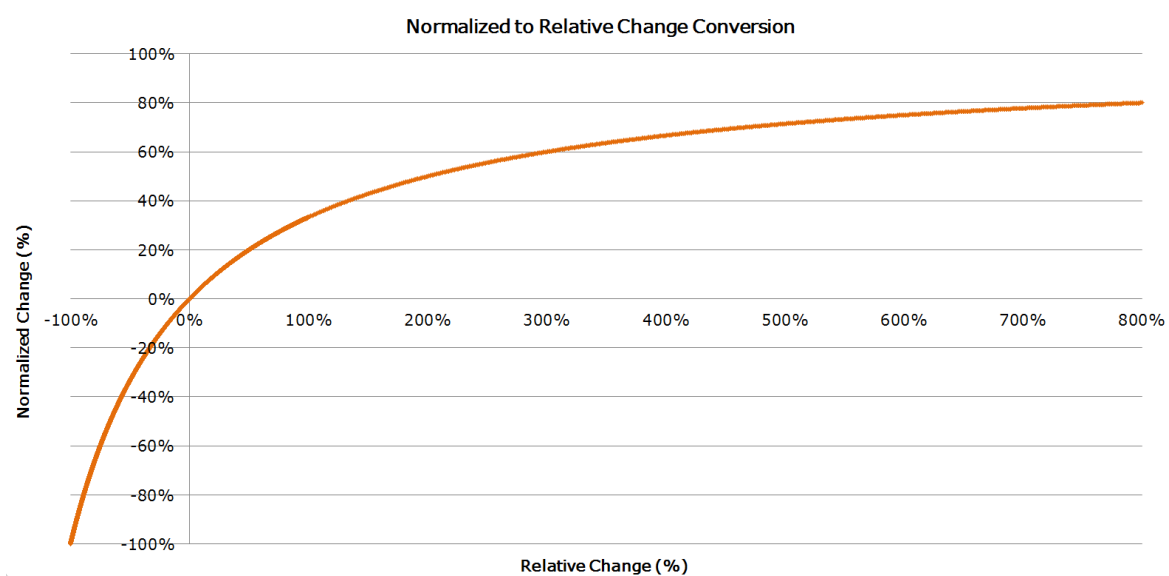


Figure S1. Normalized change (in %; values calculated by equation 1 multiplied by 100) versus relative change (%) curve.

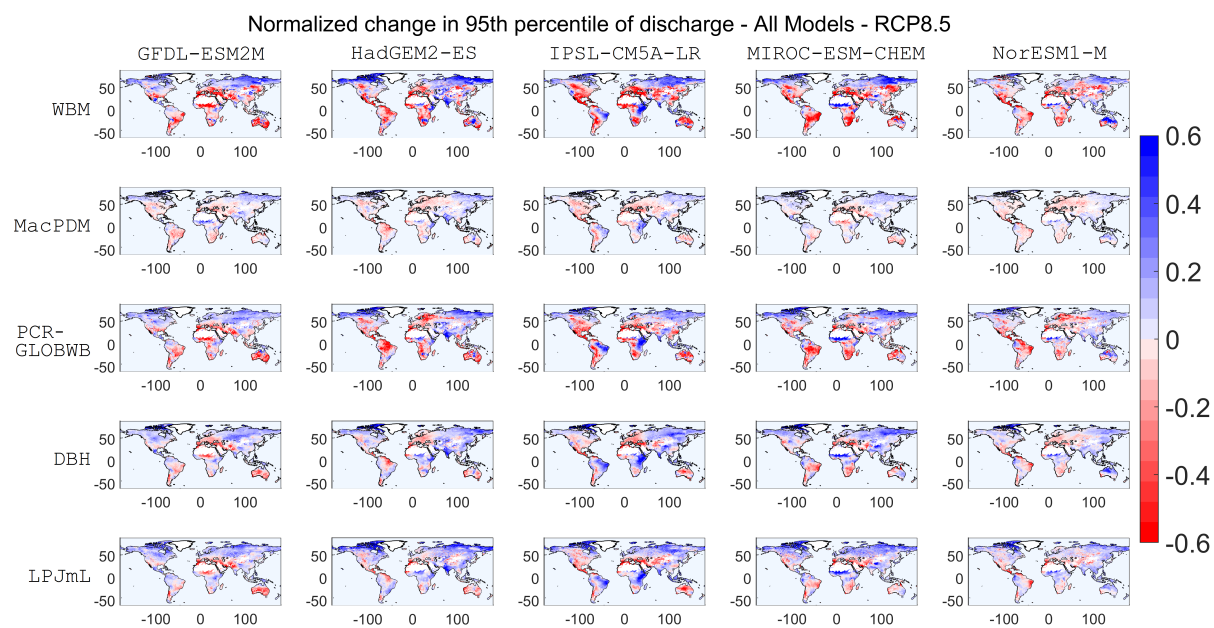


Figure S2. Normalized change in P95 for each of GHM/GCM combinations under RCP8.5 scenario.

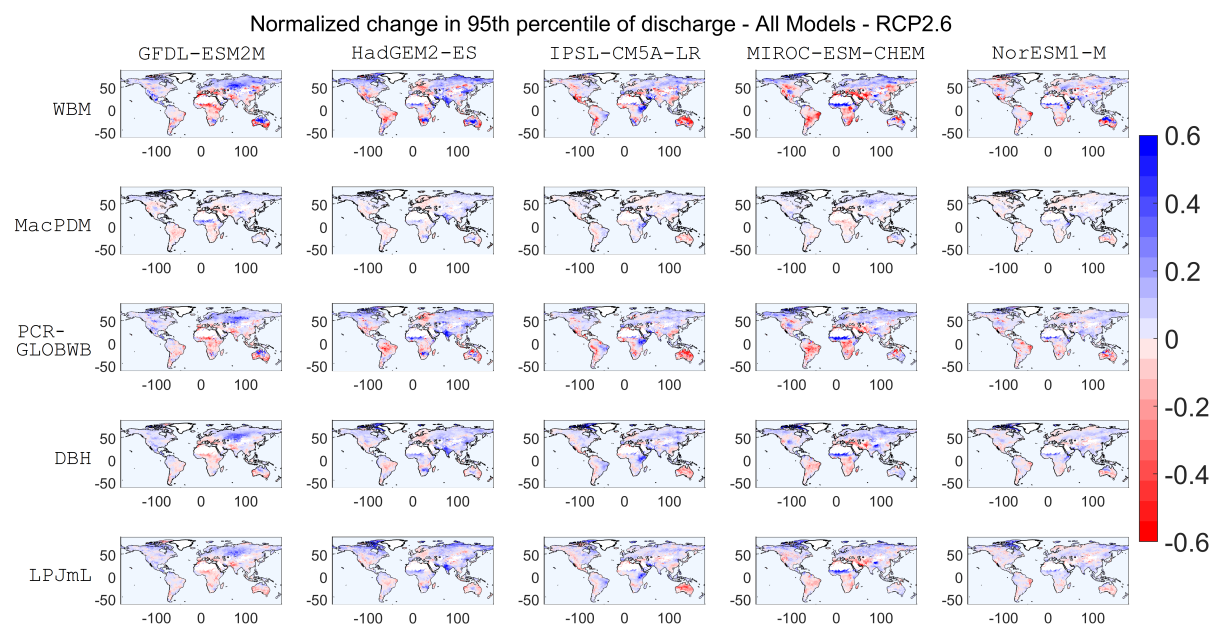


Figure S3. Normalized change in P95 for each of GHM/GCM combinations under RCP2.6 scenario.

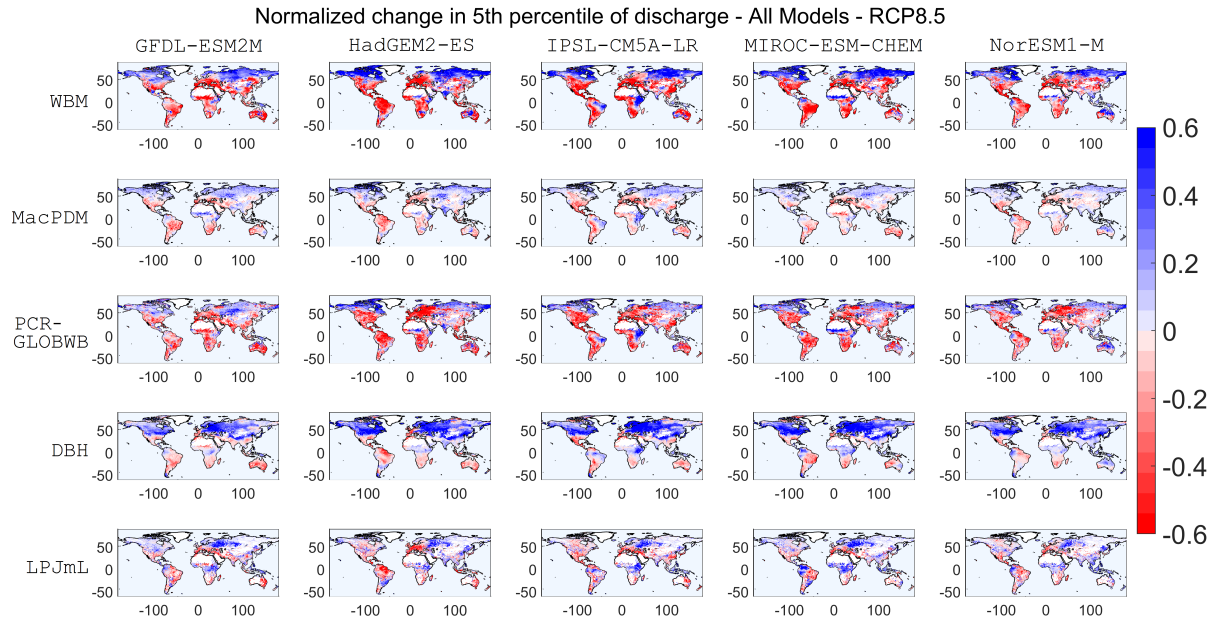


Figure S4. Normalized change in P5 for each of GHM/GCM combinations under RCP8.5 scenario.

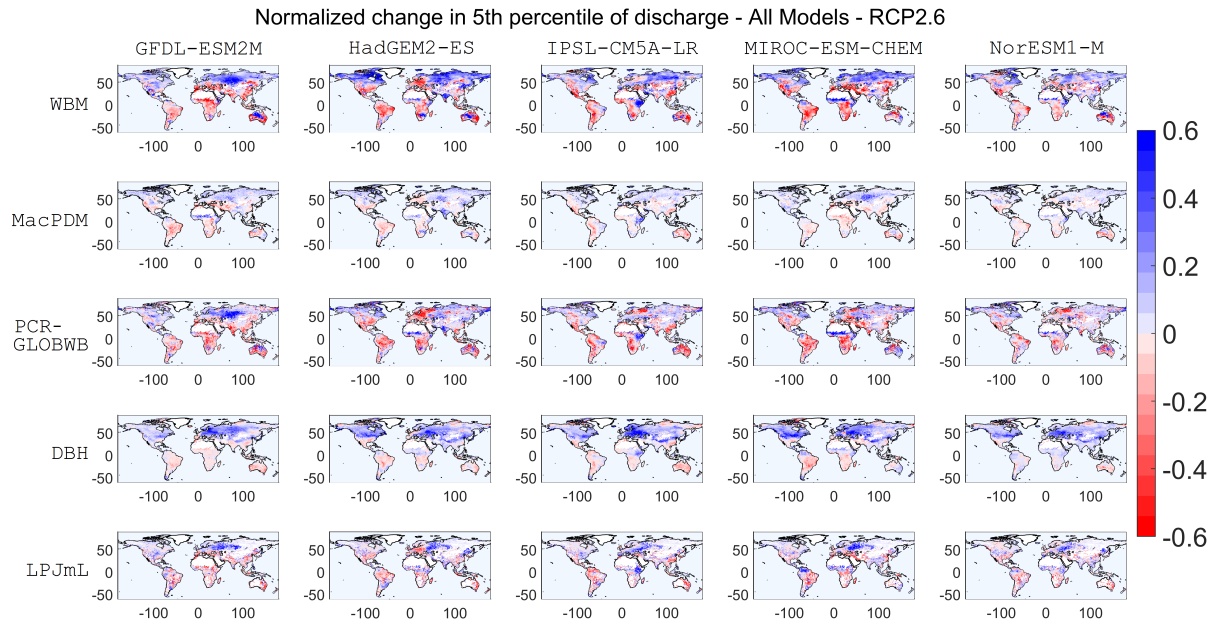


Figure S5. Normalized change in P5 for each of GHM/GCM combinations under RCP2.6 scenario.

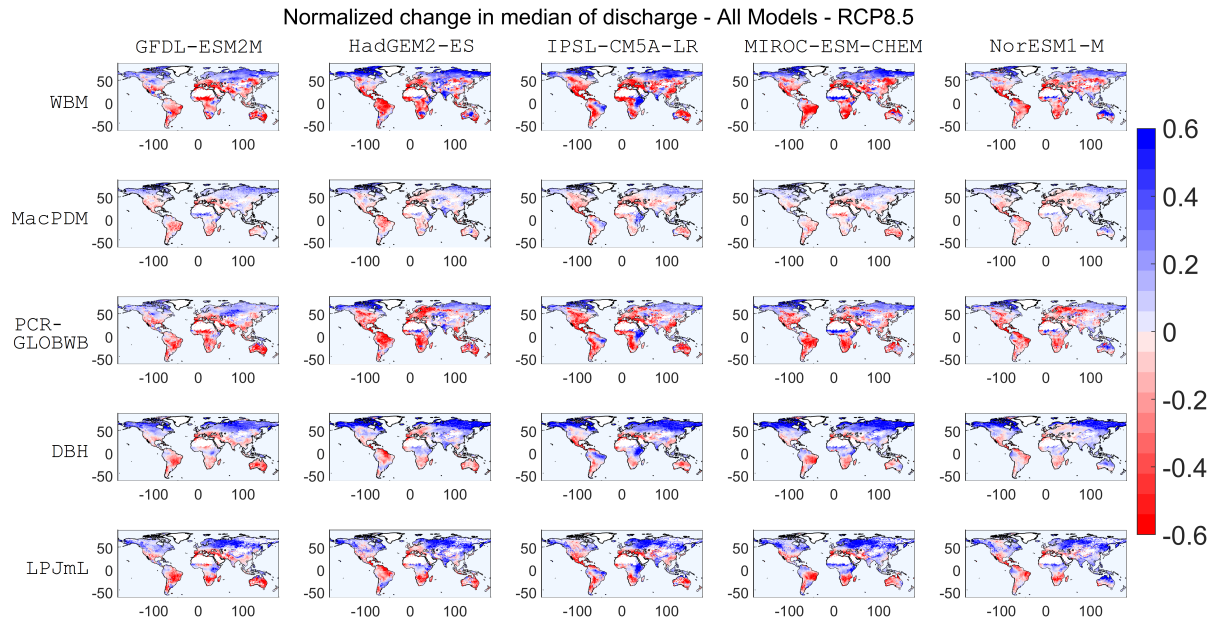


Figure S6. Normalized change in median of streamflow for each of GHM/GCM combinations under RCP8.5 scenario.

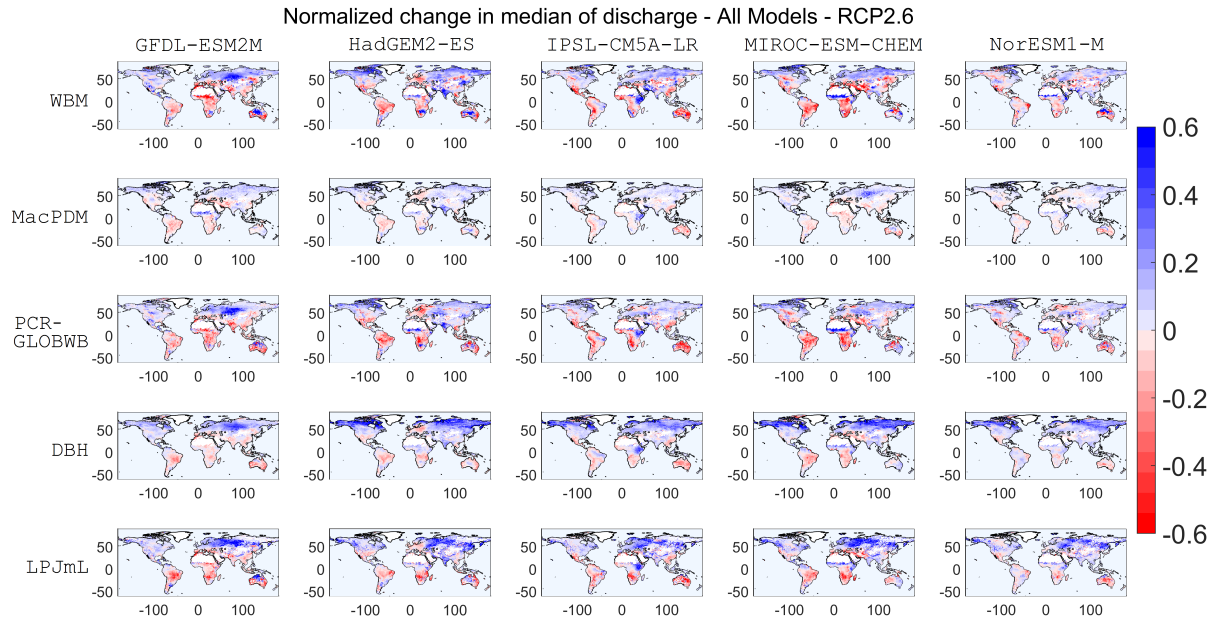


Figure S7. Normalized change in median of streamflow for each of GHM/GCM combinations under RCP2.6 scenario.