

376 **A Appendix**

377 **A.1 Proof of Theorem 1**

378 **Theorem 1.** For any sample $S = (x_1, \dots, x_m)$, the empirical Rademacher complexity of a hypothesis
 379 set \mathcal{H} is defined by $\widehat{\mathfrak{R}}_S(\mathcal{H}) = \mathbb{E}_\sigma \left[\sup_{h \in \mathcal{H}} \sum_{i=1}^m \sigma_i h(x_i) \right]$, where, σ_i s, $i \in [m]$, are independent
 380 uniformly distributed random variables taking values in $\{-1, 1\}$. The following upper bound holds
 381 for the empirical Rademacher complexity of $\mathcal{H}_{n,\lambda,q}$:

$$\widehat{\mathfrak{R}}^S(\mathcal{H}_{n,\lambda,q}) \leq \lambda \sqrt{\frac{(4n+2) \log_2(d+2) \log(m+1)}{m}},$$

382 where d is input data dimension.

383 *Proof.* For the purpose of this proof, let \mathcal{H}_n be the family of binary decision trees with leaf values
 384 $w_j \in \{-1, +1\}$. We use the regularization in the family $\mathcal{H}_{n,\lambda,q}$ and the connection to the family
 385 \mathcal{H}_n in the proof below. Additionally, let $r \geq 1$ such that $\frac{1}{r} + \frac{1}{q} = 1$, meaning that the r -norm is
 386 the dual to the q -norm. To aid the presentation in the proof, we are going to define a vector $\widehat{\sigma}$ s.t.
 387 $[\widehat{\sigma}]_j = \sum_{x_i \in \text{leaf}_j} \sigma_i$, the j -th coordinate of which contains the sum of the Rademacher variables that
 388 correspond to the sample points that fall within j -th leaf of a tree h .

$$\widehat{\mathfrak{R}}^S(\mathcal{H}_{n,\lambda,q}) = \frac{1}{m} \mathbb{E}_\sigma \left[\sup_{h \in \mathcal{H}_{n,\lambda,q}} \left[\sum_{n=1}^m \sigma_n h(x_n) \right] \right] \quad (12)$$

$$= \frac{1}{m} \mathbb{E}_\sigma \left[\sup_{h \in \mathcal{H}_{n,\lambda,q}} \left[\widehat{\sigma} \cdot \mathbf{w} \right] \right] \quad (13)$$

$$\leq \frac{1}{m} \mathbb{E}_\sigma \left[\sup_{h \in \mathcal{H}_{n,\lambda,q}} \left[\|\widehat{\sigma}\|_r \|\mathbf{w}\|_q \right] \right] \quad (14)$$

$$\leq \frac{\lambda}{m} \mathbb{E}_\sigma \left[\sup_{h \in \mathcal{H}_n} \|\widehat{\sigma}\|_r \right] \quad (15)$$

$$\leq \frac{\lambda}{m} \mathbb{E}_\sigma \left[\sup_{h \in \mathcal{H}_n} \|\widehat{\sigma}\|_1 \right] \quad (16)$$

$$= \frac{\lambda}{m} \mathbb{E}_\sigma \left[\sup_{h \in \mathcal{H}_n} \sum_{i=1}^n |[\widehat{\sigma}]_i| \right] \quad (17)$$

$$= \frac{\lambda}{m} \mathbb{E}_\sigma \left[\sup_{h \in \mathcal{H}_n} \sum_{l \in \text{leaves}(h)} \left| \sum_{i=1}^m \sigma_i 1_{\{x_i \in l\}} \right| \right] \quad (18)$$

$$\leq \frac{\lambda}{m} \mathbb{E}_\sigma \left[\sup_{h \in \mathcal{H}_n, s_l \in \{+1, -1\}} \sum_{l \in \text{leaves}(h)} s_l \sum_{i=1}^m \sigma_i 1_{\{x_i \in l\}} \right] \quad (19)$$

$$= \frac{\lambda}{m} \mathbb{E}_\sigma \left[\sup_{h \in \mathcal{H}_n, s_l \in \{+1, -1\}} \sum_{i=1}^m \sigma_i \sum_{l \in \text{leaves}(h)} s_l 1_{\{x_i \in l\}} \right] \quad (20)$$

$$\leq \lambda \sqrt{\frac{(4n+2) \log_2(d+2) \log(m+1)}{m}} \quad (21)$$

389 Where n is the number of internal nodes, and d is the input data dimension. The inequality (14) is a
 390 direct application of the Hölder's inequality for dual norms. The inequality (16) uses $\|\cdot\|_r \leq \|\cdot\|_1$.
 391 The equality (18) directly follows from the definition of $\widehat{\sigma}$. The last inequality (21) follows from the
 392 fact that the VC-dimension of binary classification trees can be bounded by $(2n+1) \log_2(d+2)$
 393 [Mohri et al. \[2012\]](#) and a direct application of Massart's lemma [Massart and Picard \[2007\]](#). \square

394 **A.2 Proof of Theorem 2**

395 **Theorem 2.** Fix $\rho > 0$. Let $\mathcal{H}_k = \mathcal{H}_{n_k, \lambda_k, q_k}$, where $(n_k), (\lambda_k)$ are sequences of constraints on
 396 the number of internal nodes n and the leaf vector norm $\|\mathbf{w}\|_q$. Define $\mathcal{F} = \text{conv}(\cup_{k=1}^K \mathcal{H}_k)$. Then,
 397 for any $\delta > 0$, with probability at least $1 - \delta$ over the draw of a sample S of size m , the following
 398 inequality holds for all $f = \sum_{t=1}^T \alpha_t h_t \in \mathcal{F}$:

$$R(f) \leq \widehat{R}_{S, \rho}(f) + \frac{4}{\rho} \sum_{t=1}^T \alpha_t \lambda_{I_t} \sqrt{\frac{(4n_{I_t} + 2) \log_2(d+2) \log(m+1)}{m}} + C(m, K),$$

399 where I_t is the index of the subclass selected at time t and $C(m, K) = O\left(\sqrt{\frac{\log(K)}{\rho^2 m}} \log \left[\frac{\rho^2 m}{\log(K)}\right]\right)$.

400 *Proof.* For this proof we are going to make use of the generalization bounds for broad families of
 401 real-valued functions given in Theorem 1 of [Cortes et al., 2014]. Adapted to our notation, it states
 402 that for any f from a family of real-valued functions \mathcal{F} that is equal to the convex hull of $\cup_{k=1}^K \mathcal{H}_k$,
 403 for any $\delta > 0$ with probability at least $1 - \delta$ over the choice of sample $S \sim \mathcal{D}^m$, the following
 404 generalization bound holds:

$$R(f) \leq \widehat{R}_{S, \rho}(f) + \frac{4}{\rho} \sum_{t=1}^T \alpha_t \mathfrak{R}_m(\mathcal{H}_t) + \frac{2}{\rho} \sqrt{\frac{\log K}{m}} + \sqrt{\left\lceil \frac{4}{\rho^2} \log \left(\frac{\rho m^2}{\log K} \right) \right\rceil \frac{\log K}{m} + \frac{\log(\frac{2}{\delta})}{2m}}.$$

405 where α_t is are the weights that represent f in the convex hull of $\cup_{k=1}^K \mathcal{H}_k$, that is $f = \sum_{t=1}^T \alpha_t h_t$
 406 s.t. $\boldsymbol{\alpha} = [\alpha_1, \dots, \alpha_T]$ is in the simplex Δ . This bound is directly applicable to the Regularized
 407 Gradient Boosting that we define, since at each boosting round, the algorithm selects a base predictor
 408 $h_t \in \mathcal{H}_t$, and multiplies it by a coefficient α_t . Thus, after T boosting rounds, we will have obtained
 409 an ensemble f such that $f = \sum_{t=1}^T \alpha_t h_t \in \text{conv}(\cup_{k=1}^K \mathcal{H}_k)$ and $\boldsymbol{\alpha}$ directly in the simplex Δ .

410 Applying the Rademacher complexity bound on the regularized families of regression trees $\mathcal{H}_{n, \lambda, q}$
 411 that we derived in Theorem 1 and noting that

$$\frac{2}{\rho} \sqrt{\frac{\log K}{m}} + \sqrt{\left\lceil \frac{4}{\rho^2} \log \left(\frac{\rho m^2}{\log K} \right) \right\rceil \frac{\log K}{m} + \frac{\log(\frac{2}{\delta})}{2m}} = O\left(\sqrt{\frac{\log(K)}{\rho^2 m}} \log \left[\frac{\rho^2 m}{\log(K)}\right]\right) \quad (22)$$

412 We obtain the expression for the bound in Theorem 2. □

413 **A.3 Proof of Lemma 3**

414 **Lemma 3.** Assume that $\Phi(y, h)$ is differentiable with respect to the second argument, and that $\frac{\partial \Phi}{\partial h}$
 415 $C_\Phi(y)$ -Lipschitz with respect to the second argument, for any fixed value y of the first argument. for
 416 all $k \in [0, K]$, define $L'_k(\boldsymbol{\alpha}) = \frac{\partial L}{\partial \alpha_k}$. Then, $L'_k(\boldsymbol{\alpha})$ is Lipschitz-continuous with the corresponding
 417 Lipschitz constants C_k bounded as follows:

$$C_k \leq \frac{1}{m} \sum_{i=1}^m h_k^2(x_i) C_\Phi(y_i). \quad (23)$$

418 *Proof.* The k -th derivative of $L(\boldsymbol{\alpha})$ is equal to (except $\alpha_k = 0$):

$$L'_k(\boldsymbol{\alpha}) = \frac{1}{m} \sum_{i=1}^m \frac{\partial \Phi}{\partial h} \left(y_i, \sum_{t=1}^T \alpha_t h_t(x_i) \right) h_k(x_i) + c_k, \quad (24)$$

419 where $c_k = \beta \lambda_k \sqrt{\frac{(4n_k+2) \log_2(d+2) \log(m+1)}{m}}$. Let \mathbf{e}_k be the k -th standard basis vector, then

$$\begin{aligned}
\left| L'_k(\boldsymbol{\alpha}) - L'_k(\boldsymbol{\alpha} + \delta \mathbf{e}_k) \right| &= \left| \frac{1}{m} \sum_{i=1}^m h_k(x_i) \left[\frac{\partial \Phi}{\partial h} \left(y_i, \sum_{t=1}^T \alpha_t h_t(x_i) \right) - \frac{\partial \Phi}{\partial h} \left(y_i, \sum_{t=1}^T \alpha_t h_t(x_i) + \delta h_k(x_i) \right) \right] \right| \\
&\leq \frac{1}{m} \sum_{i=1}^m |h_k(x_i)| \left| \frac{\partial \Phi}{\partial h} \left(y_i, \sum_{t=1}^T \alpha_t h_t(x_i) \right) - \frac{\partial \Phi}{\partial h} \left(y_i, \sum_{t=1}^T \alpha_t h_t(x_i) + \delta h_k(x_i) \right) \right| \\
&= \frac{1}{m} \sum_{i=1}^m |h_k(x_i)| \left| \frac{\partial \Phi}{\partial h} \left(y_i, f \right) - \frac{\partial \Phi}{\partial h} \left(y_i, f + \delta h_k(x_i) \right) \right| \\
&\leq \frac{1}{m} \sum_{i=1}^m |h_k(x_i)| C_\Phi(y_i) |h_k(x_i)| |\delta| \\
&= \frac{1}{m} \sum_{i=1}^m h_k^2(x_i) C_\Phi(y_i) |\delta|
\end{aligned}$$

420 Thus, $L'_k(\boldsymbol{\alpha})$ is Lipschitz-continuous with the corresponding Lipschitz constant bounded by
421 $\frac{1}{m} \sum_{i=1}^m h_k^2(x_i) C_\Phi(y_i)$.

422

□

423 A.4 Proof of Lemma 4

424 **Lemma 4.** . For each $k \in [0, K]$ let $\mathcal{H}_{n_k, \lambda_k, 2}$ be the family of regularized regression trees with
425 $\|\mathbf{w}\|_2 \leq \lambda_k$ and the number of internal nodes bounded by n_k . The regularized objective $L(\boldsymbol{\alpha})$ as
426 in Equation 7 has Lipschitz-continuous derivatives with the coordinate-wise Lipschitz constants C_k
427 bounded as follows:

$$C_k \leq \lambda_k \left[\max_{1 \leq i \leq m} C_\Phi(y_i) \right]. \quad (25)$$

428 *Proof.* For a sample S and a fixed tree h let η_l be the number of sample points falling within the leaf
429 l .

$$\begin{aligned}
C_k &\leq \frac{1}{m} \left[\max_{1 \leq i \leq m} C_\Phi(y_i) \right] \sum_{i=1}^m h_k^2(x_i) \\
&\leq \frac{1}{m} \left[\max_{1 \leq i \leq m} C_\Phi(y_i) \right] \sum_{l \in \text{leaves}(h_k)} \eta_l w_l^2 \\
&\leq \frac{1}{m} \left[\max_{1 \leq i \leq m} C_\Phi(y_i) \right] \|\mathbf{w}\|_2 \max_{l \in \text{leaves}(h_k)} \eta_l \\
&\leq \|\mathbf{w}\|_2 \left[\max_{1 \leq i \leq m} C_\Phi(y_i) \right] \\
&\leq \lambda_k \left[\max_{1 \leq i \leq m} C_\Phi(y_i) \right]
\end{aligned}$$

430 This results in the coordinate sampling distribution for the Randomized Coordinate Descent.

$$p_k = \frac{\lambda_k}{\sum_{j=1}^K \lambda_j} \quad (26)$$

431

□

Table 2: Dataset statistics

	sonar	cancer	diabetes	ocr17	ocr49	mnist17	mnist49
Examples	208	699	768	2000	2000	15170	13782
Features	60	9	8	196	196	400	400

432 **A.5 Descriptive statistics of the UCI datasets**

433 Note that mnist17 and mnist49 refer to the original 28-by-28 pixel datasets, where only two digits
434 (1,7 and 4,9 respectively) were sampled. The *cancer* dataset refers to the *breastcancer* dataset in the
435 UCI repository.