List of primitive functions

Description	In	N in	Out	N out	Comm	Param
Nominal to binary	nom	1	bin	1–4	-	Yes
Ordinal to binary	ord	1	bin	1–4	-	Yes
Linear to linear segments	lin	1	lin	1–4	-	Yes
Linear segments to binary	lin	1	bin	1–4	-	Yes
Get one column of n-matrix	bin	1-4	bin	1	-	Yes
Conjunction	bin	2-6	bin	1	Yes	-
Disjunction	bin	2–6	bin	1	Yes	-
Negate binary	bin	1	bin	1	-	-
Logarithm	lin	1	lin	1	-	-
Hyperbolic tangent sigmiod	lin	1	lin	1	-	-
Logistic sigmoid	lin	1	lin	1	-	-
Sum	lin	2-3	lin	1	Yes	-
Difference	lin	2	lin	1	No	-
Multiplication	lin,bin	2-3	lin	1	Yes	-
Division	lin	2	lin	1	No	-
Inverse	lin	1	lin	1	-	-
Polynomial transformation	lin	1	lin	1	-	Yes
Radial basis function	lin	1	lin	1	-	Yes
Monomials: $x\sqrt{x}$, etc.	lin	1	lin	1	-	-

Feature generation

There given

- the measured features $\Xi = \{\xi\}$,
- the expert-given primitive functions $G = \{g(\mathbf{b}, \xi)\}$,

$$g: \xi \mapsto x$$
;

- the generation rules: $\mathcal{G} \supset G$, where the superposition $g_k \circ g_l \in \mathcal{G}$ w.r.t. numbers and types of the input and output arguments;
- the simplification rules: g_u is not in \mathcal{G} , if there exist a rule

$$r: g_u \mapsto g_v \in \mathcal{G}$$
.

The result is

the set of the features $X = \{\mathbf{x}_1, \dots, \mathbf{x}_i, \dots, \mathbf{x}_n\}.$

The number of features exceeds the number of clients!

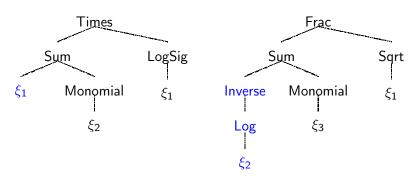
Examples of generated features

- Frac(Period of residence, Undeclared income)
- Frac(Seg(Period of employment), Term of contract)
- And(Income confirmation, Bank account)
- Times(Seg(Score hour), Frac(Seg(Period of employment), Salary))

Feature generation with symbolic regression

- Select random nodes in two features,
- 2 exchange the corresponded subtrees,
- 3 modify the function at a random node for another one from the primitive set.

Any modification must result an admissible superposition.



Feature generation with polynomials

- 1. Consider cartesian product $G \times \Xi$ of the set of non-generated variables Ξ the primitives G. Denote by a_{ι} the superpositions $g_{\nu}(\xi_{u})$
- 2. Product superpositions a_{ι} no more than P times

$$a_{\iota} = g_{\nu}(\xi_{u}),$$
 where the index $\iota = (\nu - 1)U + u$

and

$$x_j = \prod \underbrace{a_{\iota_1} \dots a_{\iota_p}}_{p \; \; \text{times}}, \quad \text{where} \;\; \iota \in \{1, \dots, UV\}, \;\; p \in \{1, \dots, P\}.$$

In the other words

$$\xi_u \xrightarrow{g_v} g_v(\xi_u) \equiv a_\iota \xrightarrow{\prod^\rho} x_j, \qquad j \in \mathcal{J}.$$

Consider the linear models as the polynomial with a monomial $a_{\iota}=g_{\nu}(\xi_{u})$

$$f(\mathbf{w},\mathbf{x}) = \sum_{\iota=1}^{UV} w_{\iota} a_{\iota} + \sum_{\iota=1}^{UV} \sum_{\kappa=1}^{UV} w_{\iota\kappa} a_{\iota} a_{\kappa} + \sum_{\iota=1}^{UV} \sum_{\kappa=1}^{UV} \sum_{\tau=1}^{UV} w_{\iota\kappa\tau} a_{\iota} a_{\kappa} a_{\tau} + \cdots$$

Set of the primitive functions *G*

Let $G = \{g_1, \dots, g_l | g = g(\mathbf{b}, \cdot, \dots, \cdot)\}$ such that there are given

- the function $g:(\mathbf{b},x)\mapsto x'$,
- its parameters **b** (the empty set is allowed),
- number of arguments v(g) of the function g and the order of the arguments (zero arguments is allowed),
- domain dom(g) and codomain cod(g).

Consider the model $f(\mathbf{w}, \mathbf{x})$ as a superposition

$$f(\mathbf{w}, \mathbf{x}) = (g_{i(1)} \circ \cdots \circ g_{i(K)})(\mathbf{x}), \text{ where } \mathbf{w} = [\mathbf{b}_{i(1)}^\mathsf{T}, \dots, \mathbf{b}_{i(K)}^\mathsf{T}]^\mathsf{T}.$$

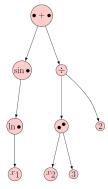
The admissible superposition f

is the superposition, which satisfies

$$cod(g_{i(k+1)}) \subseteq dom(g_{i(k)})$$
, for any $k = 1, ..., K - 1$.

The tree Γ_f corresponds to the superposition f

- The vertex V_i corresponds to the primitive function $g_{s(i)}$.
- The number of outgoing nodes from the vertex V_i equal the number of arguments of $v(g_{s(i)})$.
- The order of the outgoing nodes from the vertex V_i equals the order of the arguments of g_{s(i)}.
- The leaves of the tree Γ_f corresponds to the independent variables x_i and constants; they are treated as the primitives $g(\emptyset)$.



The tree for the superposition $\sin(\ln x_1) + \frac{x_2^3}{2}$

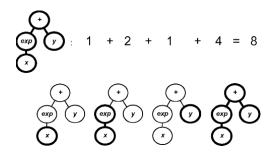
The structural density and depth

The superposition depth d(f) is

maximum depth of the tree Γ_f , number of the nodes V from the root to the most distanced leaf.

The superposition complexity C(f) is

the number of all admissible subtrees of the tree Γ_f .



Generation of nonlinear models

Given: $G = \{g_u, h_v | u \in \mathcal{U}, v \in \mathcal{V}\}$ is a set of the primitive functions of one and two arguments, $\mathbf{x} = \{x_j | j \in \mathcal{J}\}$ — independent variables.

Step 1:
$$\mathcal{F}_1 = \left\{ f_s^{(1)} \right\} = \{ g_u(x_j) \} \cup \{ h_v(x_j, x_k) \},$$

 $k \in \mathcal{J}, \ s \in \{ 1, \dots, |\mathcal{U}| \cdot |\mathcal{J}| + |\mathcal{V}| \cdot |\mathcal{J}|^2 \}.$

Step k:

(Gen) Append to \mathcal{F} the set

$$\mathcal{F}^{(k)} = \left\{ f_s^{(k)} \right\} = \left\{ g_u \left(f_{s'}^{(k-1)} \right) \right\} \cup \left\{ h_v \left(f_{s''}^{(k-1)}, f_{s'''}^{(k-1)} \right) \right\},\,$$

(Rem) which does not contain the superpositions, isomorphic to $g_u\left(f_s^{(k)}\right)$ and $h_v\left(f_s^{(k)},f_{s'}^{(k)}\right)$ form the sets $\mathcal{F}^{(k)}\ldots\mathcal{F}^{(1)}$.

Structural parameters and model selection

Exhaustive search in the set of the generalized linear models

$$\mu(y) = w_0 + \alpha_1 w_1 x_1 + \alpha_2 w_2 x_2 + \ldots + \alpha_R w_R x_R.$$

Here $\alpha \in \{0,1\}$ is the structural parameter.

Find a model defined by the set $A \subseteq \mathcal{J}$:

α_1	α_2	 $\alpha_{ \mathcal{J} }$
1	0	 0
0	1	 0
1	1	 1

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Discrete genetic algorithm for feature selection (simple ver.)

- **1** There are set of binary vectors $\{\mathbf{a}_1, \dots, \mathbf{a}_P\}$, $\mathbf{a} \in \{0, 1\}^n$;
- 2 get two vectors $\mathbf{a}_p, \mathbf{a}_q, p, q \in \{1, \dots, P\}$;
- **3** chose random number $\nu \in \{1, \dots, n-1\}$;
- 4 split both vectors and change their parts:

$$[a_{p,1},\ldots,a_{p,\nu},a_{q,\nu+1},\ldots,a_{q,n}] \rightarrow \mathbf{a'}_p,$$

$$[a_{q,1},\ldots,a_{q,\nu},a_{p,\nu+1},\ldots,a_{p,n}]\to {\bf a'}_q;$$

- **5** choose random numbers $\eta_1, \ldots, \eta_Q \in \{1, \ldots, n\}$;
- **6** invert positions η_1, \ldots, η_Q of the vectors $\mathbf{a'}_p, \mathbf{a'}_q$;
- $\mathbf{7}$ repeat items 2-6 P/2 times;
- 8 evaluate the obtained models.

Repeat R times; here P, Q, R are the parameters of the algorithm and n is the number of the corresponding model features.

Discrete genetic algorithm for grouping

- **1** There are set of binary vectors $\{\mathbf{a}_1,\ldots,\mathbf{a}_P\}$, $\mathbf{a}\in\{1,\ldots,k\}^n$;
- 2 get two vectors $\mathbf{a}_p, \mathbf{a}_q, p, q \in \{1, \dots, P\}$;
- **3** chose random number $\nu \in \{1, \dots, n-1\}$;
- 4 split both vectors and change their parts:

$$[a_{p,1},\ldots,a_{p,\nu},a_{q,\nu+1},\ldots,a_{q,n}] o \mathbf{a'}_p,$$

 $[a_{q,1},\ldots,a_{q,\nu},a_{p,\nu+1},\ldots,a_{p,n}] o \mathbf{a'}_q;$

- **5** choose random numbers $\eta_1, \ldots, \eta_Q \in \{1, \ldots, n\}$;
- **6** replace values in positions η_1, \ldots, η_Q of the vectors $\mathbf{a'}_p, \mathbf{a'}_q$ for random values from $\{1, \ldots, k\}$;
- \bigcirc repeat items 2-6 P/2 times;
- 8 evaluate the obtained models.

Repeat R times; here P, Q, R are the parameters of the algorithm and k is desired number of categories.