# Data Dictionary 

# for <br> Vegetation Resources Inventory and <br> National Forest Inventory <br> Timber Data 

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## Data Dictionary for Vegetation Resources Inventory and National Inventory Summary Files

The VRI compiler summarizes tree data at the cluster level and the cluster/species level. Compiled results are stored in the VRI database in files named SMY_C (cluster level) and SMY_CS (cluster/species level). The following table describes fields stored in these files. There are a number of fields that occur on both files, the description of the files has been combined. These files SMY_NC and SMY_NCS are identical to SMY_C and SMY_CS with 4 additional fields. These fields are also applicable to the Net Volume Adjustment Factor (NVAF) application. In some files, these fields may be missing and would indicate that no NVAF is available for the data.

| Identifier | mation |  |
| :---: | :---: | :---: |
|  | Found in files | Variable label, and description. |
| Clstr_Id | Smy_nc \& smy_ncs | Project Id - Sample \# - Type Code <br> Concatenation of the Project identifier, Sample number and type code. |
| Proj_id | Smy_nc \& smy_ncs | Project identifier <br> The province has been divided into a number of inventory units and a project code is provided. Project identifier also included in Clstr_id above. |
| Samp_no | Smy_nc \& smy_ncs | Sample Number <br> Each ground sample cluster is assigned a number |
| Type_cd | Smy_nc \& smy_ncs | Type code <br> Is a 3 character code indicating the type of sampling, intent of sampling and measurement \#. It could be an audit, NVAF, NFI/CMI sample, etc. (see VRI field sampling procedure manual for all codes) |
| Prj_grp | Smy_nc \& smy_ncs | Project Grouping <br> A grouping of projects used for regression analysis. |
| No_plots | Smy_nc \& smy_ncs | \# of plots in the sample <br> This ranges from 1 to 5 . There can be an Integrated plot at the centre(IPC) and up to 4 auxiliary plots. |
| Blowup | Smy_nc \& smy_ncs | Blowup Factor <br> A factor which applies to an individual tree on a per hectare basis. For variable area plots (V) - Basal Area Factor (BAF) of plot For fixed area plots (F) - 1/Plot area |
| Bgc_zone, Bgc_sbzn, bgc_var | Smy_nc \& smy_ncs | Bec Zone, subzone, and variant. <br> These are biogeoclimatic (BEC) designations as per "Ecosystems of BC " (Meidinger and Pojar, 1991) They are used to compute volume using Kozak's taper equation and to check for certain adjustment factors, such as the interior wetbelt. |
| Fiz | Smy_nc \& smy_ncs | Forest Inventory zone: <br> Twelve forest inventory zones covering the province, resembling BEC zones. Previously used for Kozak's volume equation and loss factors |
| Samp_Typ | Smy_nc \& smy_ncs | Sample Type F/V <br> Indicates if the sample is fixed area or variable radius (prism or relaskope). |


| Species | smy_ncs | Species <br> Tree species, used at the species summary level. Genus and species are <br> collected |
| :--- | :--- | :--- |
| Sp0 | Smy_ncs | Species grouping - <br> Grouping of tree species, used for Kozak's volume equation as well as <br> Util |
|  |  <br> smy_ncs | fitting regression models. <br> Utilization Level - |
| The minimum tree DBH included in calculations. The files store <br> values compiled to $4 \mathrm{~cm}, 7.5 \mathrm{~cm}, 12.5 \mathrm{~cm}, 17.5 \mathrm{~cm}$ and $22.5 \mathrm{~cm} . ~ I f ~$ <br> required, other utilization levels can be computed. |  |  |

## Age and height related data - Based on data collected on the "Site tree data - TS card " and site tree info from "Auxiliary Plot card - TA"

| Variable | Found in |
| :--- | :--- |
| name | files |
| Ab_m_tls | Smy_nc |
| Ageb_tls | smy_ncs |


| Ab_m_txo | Smy_nc <br> smy_ncs | Mean Age @Breast Height for T,L,S,X $\mathbf{X}$ and $\mathbf{O}$ trees <br> The calculation is as above, but the trees included in this average are <br> tree types L, S, T, extra(X)and other(O) |
| :--- | :--- | :--- |
| At_m_tls | Smy_nc <br> smy_ncs | Mean Total Age for T,L and S trees <br> The calculation is as above, but the age may be derived or taken <br> directly from input data.. Breast height age is usually captured. Site <br> index equations (MOF, Research Br, Site Tools) are used to convert <br> from breast height age to total age. Trees included in this average are <br> only type L, S and T. |
| Aget_tls |  |  |


| $\begin{aligned} & \text { Ht_m_txo } \\ & \text { Ht_txo } \end{aligned}$ | Smy nc | Height Mean for T,L,S,X and O trees |
| :---: | :---: | :---: |
|  | smy_ncs | The calculation is as above, but the trees included in this average are types L, S, T, X and O. |
| Ht_Rnd | Smy_nc | Height Random Tree |
|  |  | Height of "R" (Random) tree as taken from the sample data |
| Ht_Top | smy_nc | Height "Top Height" Tree |
|  |  | Height of Top Height (T) tree as taken from the sample data. For data prior to 1999 field season cards, there were multiple T trees. This value is defined as the Height of the T tree on the IPC, of which there should be only one. |
| N_ag_tls | Smy_nc \& | \# of ages used for TLS trees |
|  | Smy_ncs | Number of ages used in the calculation of both breast height and total age for T, L, and S trees. With 1999 field season cards, an age can be designated as representative or not representative of the stand. Prior to 1999 , all ages were deemed to be representative. |
| N_ag_txo | Smy_nc \& | \# of ages used for TLSXO trees. |
|  | Smy_ncs | As above, but for all site tree types - T, L, S, X, and O |
| N_ht_tls | Smy_nc \& | \# of heights used for TLS trees |
|  | Smy_ncs | Number of heights used in the calculation of mean height for species and tree types T, L, and S trees. With 1999 field season cards, a height can be designated as representative or not representative of the stand. Prior to this, all ages were deemed to be representative. Note that a tree may be representative for height and not age and vice versa. |
| N_ht_txo | Smy_nc \& | \# of heights used for TLSXO trees. |
|  | Smy_ncs | As above, but for all site tree types - T, L, S, X, and O |
| Si_c_tls | Smy_ncs | Site index Calc for TLS trees. |
|  |  | This site index is calculated using the average breast height age (Ageb_tls) and the mean height (Ht_tls) with Site tools (as provided by Research Branch). It is possible to have a calculated site index but not a mean site index. Site index is an expression of the forest site quality of a stand, at a specified age based on site height. |
| Si_c_txo | Smy_ncs | Site index Calc for TLSX and 0 trees. <br> This site index is calculated using the average breast height age (Ageb_txo) and the mean height (Ht_txo) using Site tools (as provided by Research Branch). It is possible to have a calculated site index but not a mean site index. |
| Si_m_tls | Smy_ncs | Site index Mean for TLS trees. |
|  |  | This site index is calculated as the mean of the site indexes for trees where there is a representative height and age on the sample data. This is only for trees with types T, L and S. |
| Si_m_txo | Smy_ncs | Site index Mean for TLS trees. |
|  |  | This site index is calculated as the mean of the site indexes for trees where there is a representative height and age on the sample data. This is only for trees with types T, L, S, X and O trees. |
| Sp_Rnd | Smy_nc | Species Of Random Tree |
|  |  | Species of the randomly selected "R" tree as taken from sample data |
| Sp_Top | Smy_nc | Species Of Top Height Tree |
|  |  | Species of "T" tree as taken from sample data. For the data prior to 1999 field season, this is the T tree on the IPC. Top height tree is the tree with the largest diameter in the IPC within a 5.64 radius plot, providing it is suitable. See VRI sampling procedures for definition of suitable. |


| Volume a TD", Tre | sal are sureme | ated data - Based on data collected on the "Tree Details and live standing trees from "Auxiliary Plot card - TA" |
| :---: | :---: | :---: |
| Variable name | Found in files | Variable label, and description. |
| Ba_Ha | Smy_nc \& smy_ncs | Basal Area Per ha (Live) <br> Basal Area/Ha = Sum Of (Tree Basal Area * Tree Per Hectare Factor*Plot Wt.)/no. of plots. This is for live trees. (same as BAF * no. of trees, with appropriate plot weights). Basal area is the crosssection of tree stems at DBH, including bark, measured over 1 ha |
| Ba_Had | Smy_nc \& smy_ncs | Basal Area Per ha (Dead) <br> As above, for dead trees. (Dead trees are tallied on the IPC only; no. of plots =1), except for NVAF samples (type_cd = 'Nxx'), where dead trees can be on all auxiliary plots. |
| Ba_Pc | smy_ncs | \% Species By B.A/Ha Live <br> The species basal area/ha live expressed as a percentage of the total live basal area / ha. |
| Dha_Mer | smy_ncs | \$/Ha (Mer) - Live |
| Dht_mer | smy_nc | A $\$$ value based on a static dollar value assigned to species and log grades values, which is then used to calculate a possible dollar value for the merchantable volume for a species in a sample. The table of values was created about 1999 and has not been updated, nor are there plans to keep it updated. Merchantable volume of a tree is that which has attained a size, quality and/or volume sufficient to make it suitable for harvesting (or Whole stem volume less volume of top and stump) |
| Dha_Merd | smy_ncs | \$/Ha (Mer) - Dead |
| Dht_merd | smy_nc | As above for dead trees |
| Ht_Mean1 | Smy_nc | Height Wt. Mean All Full Measured Trees <br> Mean height based on a weighted ${ }^{1}$ mean of live standing full measured trees - includes broken top trees |
| Ht_Mean2 | Smy_nc | Height Wt. Mean Full Top Measured Trees <br> Mean height based on a weighted mean of live standing full measured trees which are not broken top |
| Ht_Mnall | Smy_nc | Height Wt. Mean All Full Measured Trees <br> Mean height based on a weighted mean of all (live and dead) standing full measured trees - includes broken top trees |
| Qmd | Smy_nc \& smy_ncs | Quad. Mean Diameter (Live) <br> Sqrt [Sum ( Dbh**2 * Phf_Tree) / Sum (Phf_Tree)] for live trees The quadratic mean gives additional weight to larger trees. |
| Qmdd | Smy_nc \& smy ncs | Quad. Mean Diameter (Dead) as above for dead trees. |
| Spb_cpet | Smy_nc | Species Composition by Basal area <br> Is a list of the top 5 species by $\%$ of basal area ranked highest to lowest |
| Stems_Ha | Smy_nc \& smy_ncs | $\begin{aligned} & \text { Stems/Ha (Live) } \\ & \text { Stems Per Hectare }=[\text { Sum (Phf_Tree of live trees) }] / \text { No Of Plots, } \end{aligned}$ |

[^0]calculated at 5 standard utilization levels. This is often referred to as density or stand density.

| Stems_Hd |  <br> smy_ncs |
| :--- | :--- | :--- | | Stems/Ha (Dead) |
| :--- |
| as above for dead trees. |

The volumes below are listed as much as possible in decreasing values.

| Vha_Wsv | smy_ncs |  |
| :---: | :---: | :---: |
| Vht_wsv | smy_nc | Whole stem volume/ha of living trees by species. |
|  |  | Calc. Vha_wsv = sum (vol_wsv* basal area of tree) trees in group Vht_wsv = sum of species vha_wsv. Volume of the main stem, including stump and top as well as defective and decayed wood of the tree. This is often referred to as "gross volume." |
| Vha_Mer | smy_ncs | Whole stem volume less top and stump / ha (live) |
| Vht_mer | smy_nc | This is whole stem volume less volume of top (set at 10 cm unless otherwise specified), and volume of stump (stump height is 30 cm ). Vha mer is species level, vht mer is a summary by sample |
| Vha_Net | smy_ncs | Whole stem Vol./Ha less Cruiser Decay (live) |
| Vht_net | smy_nc | This is whole stem volume less cruiser called decay. Vha_net is species level, vht net is a summary by sample |
| Vha_Netm | smy_ncs | Whole stem Vol./Ha less Top, Stump \&Cruiser Decay (live) |
| Vht_netm | smy_nc | This is whole stem volume less cruiser called decay, volume of top (set at 10 cm unless otherwise specified), and volume of stump. |
| Vha_Nw2 | smy_ncs | Whole stem vol/ha less Top, Stump, Cruiser Decay and Waste (live) |
| Vht_nw2 | smy_nc | This is whole stem volume less cruiser called decay, sections of tree where the cruiser decay is $>50 \%$, volume of top (set at 10 cm unless otherwise specified), and volume of stump. If more than $50 \%$ of the tree is waste then this volume is 0 . |
| Vha_Nwb | smy_ncs | Vol./Ha Live -Top, Stump, Cruiser Decay, Waste and Breakage |
| Vht_nwb | smy_nc | Species level net Merch. per hectare volume. This is whole stem volume less cruiser called decay, sections of tree where the cruiser decay is > $50 \%, \%$ breakage as per Biogeclimatic(BEC) based loss factor tables, volume of top (set at 10 cm unless otherwise specified), and volume of stump. |
| Nvl_nwb | Smy_nc \& smy_ncs | Nvl_nwb (smy_ncs) = Vha_nwb * nvaf value by strata for sample nvl_nwb (smy_nc) = sum of vha_nwb values by utilization and sample \# <br> Whole stem volume less cruiser called decay, volume of waste, top and stump (as defined above) * Net volume adjustment factor by strata. NVAF is usually applied by strata - which could be species groupings, age, or geographic area. |
| Vha_Wsvd | smy_ncs | Total Vol./Ha (Ws.Dead) |
| Vht_wsvd | smy_nc | See above vha_wsv for definition |
| Vha_Merd | smy_ncs | Vol./Ha Less Top And Stump (dead) |
| Vht_merd | smy_nc | See above vha_mer / vht_mer for exact definition |
| Vha_Netd | smy_ncs | Vol./Ha - Cruiser Decay (dead) |
| Vht_netd | smy_nc | See above vha_net / vht_net for exact definition |


| Vha_Ntmd <br> Vht_ntmd | $\begin{aligned} & \text { smy_ncs } \\ & \text { smy_nc } \end{aligned}$ | Vol./Ha -Top, Stump \&Cruiser Decay (dead) See above vha_netm / vht_netm for exact definition |
| :---: | :---: | :---: |
| Vha_Nw2d | smy_ncs | Vol./Ha -Top, Stump, Cruiser Decay and Waste (dead) |
| Vht_nw2d | smy_nc | See above vha_nw2 / vht_nw2 for exact definition. |
| Vha_Nwbd <br> Vht_nwbd | $\begin{aligned} & \text { smy_ncs } \\ & \text { smy_nc } \end{aligned}$ | Vol./Ha dead -Top, Stump, Cruiser Decay, Waste and Breakage (dead) |
|  |  | See above vha_nwb / vht_nwbd for exact definition |
| Nvl_nwbd | $\begin{aligned} & \text { Smy_nc \& } \\ & \text { smy_ncs } \end{aligned}$ | Net dead m3/ha volume (vha_nwbd, or vht_nwbd) with Net Volume Adjustment Factor (NVAF )dead factor applied. |
| Vpc_Wsv | smy_ncs | $\% \text { Vol./Ha (Ws.Live) }$ <br> For a species live whole stem volume as a $\%$ of total sample whole stem volume calc: (vha_wsv/vht_wsv) |
| Vpc_Mer | smy_ncs | \%Vol./Ha (Mer) <br> For a species Merch. volume as a $\%$ of total sample whole stem merch. volume <br> Calc: (vha_mer/vht_mer) |
| Vpc_Netm | smy_ncs | \%Vol./Ha (Net.Ws) <br> For a species net volume as a \% of total sample whole stem less volume Calc (vha_netm/vht_netm) |
| Vpc_Wsd | smy_ncs | \%Vol./Ha (Ws.Dead) <br> For a species dead whole stem volume as a $\%$ of total sample dead whole stem volume calc: (vha_wsvd/vht_wsvd) |
| Nvafwd_d | Smy_ncs | Nvaf_factor dead volume <br> The nvaf factor which is applied to dead volume. The factor may be stratified in many ways (ie age, species etc).(Not always calculated). |
| Nvafwd_l | Smy_ncs | Nvaf factor live volume <br> As above |
| Strata_info | $\begin{aligned} & \text { Smy_nc \& } \\ & \text { smy_ncs } \end{aligned}$ | Free format text field identifying the Nvaf strata used for this particular sample - |

## Volumes are as follows:

_WSV/WSVD = Whole Stem(Gross whole stem) i.e. the entire tree
_MER/MERD = Merch: whole stem less top (usually 10 cm ) and stump (usually 30 cm)

- Volumes Net of Cruiser called decay (from the field cards)
_NET/NETD = Net: whole stem volume less cruiser called decay
_NETM/NTMD = Net Merch. Whole stem volume less top, stump and cruiser called decay
_NW2/NW2D = Net merch. and waste: whole stem less top, stump, cruiser called decay and any section of the tree where the decay is $>50 \%$. If more than $50 \%$ of the tree has been removed due to waste, then the volume is 0 . Waste is the unusable volume due to its proximity to decay exceeding $50 \%$ of the volume of any section or the whole tree.
$N W B / N W B D=$ Net merch. waste and breakage: whole stem volume less top, stump, cruiser called decay, waste as described above, and breakage per the loss factor tables using BEC based breakage to determine the \% deducted. Breakage is volume of wood that becomes non merchantable when stems break into pieces too small to be handled economically under current logging methods.


## The following prefixes indicate:

VHA $=$ a $\mathrm{m} 3 /$ ha volume which is summarized to the sample and species level
VHT $=$ a $\mathrm{m} 3 /$ ha volume which is summarized to the sample level
$\mathbf{N V A F}_{-}=\mathrm{a} \mathrm{m} 3 /$ ha volume to which an NVAF factor has been applied: (currently NVL_NWB, NVL_NWBD)

A suffix of $\mathbf{D}$ generally indicates Dead volume.


[^0]:    ${ }^{1}$ Weight refers to the method used to select the tree. For fixed radius plots the probability a tree is selected from the polygon, is proportional to frequency (PPF sampling) of occurrence. For variable radius plots the probability of a tree being selected is proportional to the size (PPS sampling) of the tree.

